WORLD INTELLECTUAL PROPERTY ORGANIZATE International Bureau



INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification 5: WO 94/12162 (11) International Publication Number: A1 A61K 31/135, 37/02 9 June 1994 (09.06.94) (43) International Publication Date:

(21) International Application Number:

PCT/FI93/00514

(22) International Filing Date: .

1 December 1993 (01.12.93)

(30) Priority Data:

988,427

1 December 1992 (01.12.92)

(71)(72) Applicants and Inventors: WÄRRI, Anni, Maija [FI/FI]; Juhaninkuja 2 as. 2, FIN-21420 Lieto (FI). ALANEN-KURKI, Leena, Maria [FI/FI]; Piispankatu 6 E 31, FIN-20500 Turku (FI). AUVINEN, Petri, Olli, Viljami [FI/DE]; Kirschgartenstrasse 5, Rohrbach, D-69126 Heidelberg (DE). JAAKKOLA, Panu, Mikko [FI/FI]; Kellonsoittajankatu 13 B 20, FIN-20500 Turku (FI). JALKANEN, Markku, Tapani [FI/FI]; Rauvolantie, FIN-20760 Piispanristi (FI). LEPPÄ, Sirpa, Marianne [FI/FI]; Valkintie 14 as 6, FIN-20660 Littoinen (FI). MALI, Markku, Sakari [FI/FI]; Inkereentie 176, FIN-24280 Salo (FI). VIHINEN, Tapani, Artero [FI/FI]; Kaskenkatu 11 C 54, FIN-20700 Turku (FI).

(74) Agent: ORION CORPORATION; Orion-Farmos Pharmaceuticals, Patent Department, P.O. Box 65, FIN-02101 Espoo (81) Designated States: AT, AU, BY, CA, CH, CZ, DE, DK, ES, FI, GB, HU, JP, KR, LU, LV, NL, NO, NZ, PL, PT, RU, SE, SK, UA, US, European patent (AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE).

Published

With international search report.

Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.

(54) Title: SYNDECAN STIMULATION OF CELLULAR DIFFERENTIATION

(57) Abstract

Methods are provided for altering levels of syndecan within a cell. The methods include enhancing syndecan expression via administration of growth factors, preventing suppression of syndecan expression via administration of anti-steroid agents, and altering syndecan biochemistry within the cell. The methods are used to induce or maintain cellular differentiation, and to decrease the growth of malignant cells. Application of the methods to the treatment of patients, including humans, is provided.

BEST AVAILABLE COPY

FOR THE PURPOSES OF INFORMATION ONLY

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

					•
AT	Austria	GB	United Kingdom	MOR	Mauritania
ΑÜ	Australia	GE	Georgia	MW	Malawi
BB	Barbados	GN	Guinea	NE	Niger
BE	Belgium .	GR	Greece	NL	Netherlands
BF	Burkina Faso	HU	Hungary	NO	Norway
BG	Bulgaria	IDE	Ireland	NZ	New Zealand
BJ	Benin	īT	Italy	PL	Poland
BR	Brazil	JP	Japan	PT	Portugal .
BY	Belarus	KE	Kenya	RO	Romania
CA	Canada	KG	Kyrgystan	RU	Russian Federation
CF	Central African Republic	KP	Democratic People's Republic	SD	Sudan
CG	Congo		of Korea	SE	Sweden
CH	Switzerland	KR	Republic of Korea	SI	Slovenia
CI	Côte d'Ivoire	KZ	Kazakhstan	SK	Slovakia
CM	Carneroon	LI	Liechtenstein	SN	Senegal
CN	China	ĹK	Sri Lanka	TD	Chad
cs	Czechoslovakia	LU	Luxembourg	TG	Togo
cz	Czech Republic	LV	Latvia	TJ	Tajikistan
DE	Germany	MC	Monaco	TT	Trinidad and Tobago
DK	Denmark .	2.00	Republic of Moldova	UA	Ukraine
ES	Saeta	MG	Madagascar	US	United States of America
FI	Finland	MIL '	Mali	UZ.	Uzbekistan
FR	Prance	MN	Mongolia	VN	Vict Nam
GA	Gabon	1721			
UΛ	Central				

SYNDECAN STIMULATION OF CELLULAR DIFFERENTIATION

FIELD OF THE INVENTION

This invention is in the field of cancer biology and therapy. Specifically, the invention is to a method for altering the differentiated state of a cell by altering syndecan expression. The method allows for the normalization of the growth rate and differentiation state of malignant cells, such method being based on the stimulation of syndecan expression in malignant cancer cells. Reexpression of syndecan in such malignant cells promotes their normal differentiated phenotype and prevents their tumor formation. This method may also be applied to normal cells to increase their differentiation, and therefore support the maintenance of cells, e.g. keratin production to prevent baldness.

BACKGROUND OF THE INVENTION

It is becoming more evident that cell surface proteoglycans play an important role in the regulation of cell behavior (Ruoslahti et al., Cell 64:867-869 (1991)). Through their covalently bound glycosaminoglycan side chains, such proteoglycans can bind various extracellular effector molecules (Jalkanen, et al., in Receptors for Extracellular Matrix, J. MacDonald & R. Mecham, Editors, Academic Press, San Diego, pp. 1-37 (1991)). One central challenge in proteoglycan biology is the need to understand what directive a cell receives by binding different effector molecules via the cell surface proteoglycans. It is further imperative to investigate what kind of intracellular response such binding activates, thereby leading to altered behavior of the cell. The way to approach these questions is to create biological models which are dependent on the expression of any given proteoglycans.

Syndecan is the best characterized cell surface proteoglycan (Saunders et al., J. Cell Biol. 108:1547-1556 (1989); Mali et al., J. Biol. Chem. 265:6884-6889 (1990)). It was originally isolated from mouse mammary epithelial (NMuMG) cells as a hybrid proteoglycan containing both heparin sulfate and chondroitin sulfate glycosaminoglycan side chains (Rapraeger et al., J. Biol. Chem. 260:11046-11052 (1985)). Recent studies have revealed its expression, not only on epithelial cells but also on differentiating fibroblasts of developing tooth (Thesleff et al., Dev. Biol. 129:565-572 (1988); Vainio et al., J. Cell Biol.

108:1945-1964 (1989)), on endothelial cells of sprouting capillaries (Elenius et al., J. Cell Biol. 114:585-596 (1991)) and on the surface of lymphocyte subpopulations (Sanderson et al., Cell Regul. 1:27-35 (1989)) intimating that its function can vary on the surfaces of different cells. Syndecan belongs to a family of proteoglycans with conserved plasma membrane and cytoplasmic domains but with dissimilar ectodomains (Mali et al., J. Biol. Chem. 265:6884-6889 (1990)). The conserved structure of syndecan suggests that it could participate in signal transduction through the plasma membrane.

Syndecan binds several extracellular effector molecules but this binding is selective. For example, syndecan binds interstitial collagens and fibronectin but does not bind to vitronectin or laminin (Koda et al., J. Biol. Chem. 260:8156-8162 (1985)); Saunders et al., J. Cell Biol. 106:423-430 (1988); Elenius et al., J. Biol. Chem. 265:17837-17843 (1990)). Moreover, syndecan isolated from tooth mesenchyme has revealed selective binding to tenascin not observed for syndecan from NMuMG cells (Salmivirta et al., J. Biol. Chem. 266:7733-7739 (1991)). This suggests variation in syndecan glycosylation that results in the selective binding properties for syndecan. Polymorphism of syndecan glycosylation has also been observed between simple and stratified epithelia (Sanderson et al., Proc. Natl. Acad. Sci. USA 85:9562-9566 (1988)); but whether these changes also reflect altered ligand recognition by syndecan remains unknown. Syndecan also binds growth factors, such as basic fibroblast growth factor (bFGF) (Kiefer et al., Proc. Natl. Acad. Sci. USA 87:6985-6989 (1990); Elenius et al., J. Biol. Chem. 267:6435-6441 (1992)). Very recently, Yayon and coworkers (Yayon et al., Cell 64:841-848 (1991)) and Rapraeger and co-workers (Rapraeger et al., Science 252:1705-1708 (1991)) have shown that heparin-like molecules are required for the binding of bFGF to its high affinity receptor, indicating that syndecan-like molecules can also modulate the growth factor response. The fact that cell surface proteoglycans can bind both growth factors and matrix components could theoretically imply a role in regulating, both temporally (timing of expression) and spatially (precise localization), growth promotion by immobilizing these effector molecules to the vicinity of cell-matrix interactions. This is supported by the intriguing pattern of syndecan expression in the development that follows morphogenetic rather than histological patterns (Thesleff et al., Dev. Biol. 129:565-572 (1988); Vainio et al., J. Cell Biol. 108:1945-1954 (1989) and Vainio et al., Dev. Biol. 134:382-391 (1989)), and moreover, that syndecan expression is localized to sites of active proliferation (Elenius et al., J. Cell Biol. 114:585-596 (1991) and Vainio et al., Dev. Biol. 147:322-333 (1991)).

In simple epithelium, syndecan is polarized to baso-lateral surfaces where it co-localizes with actin rich cytofilaments (Rapraeger et al., J. Cell Biol. 103:3683-2696 (1986)). Upon rounding, syndecan is shed from the cell surface by proteolytic cleavage of the core protein at the cell surface, a process which separates the matrix binding ectodomain from the membrane domain (Jalkanen et al., J. Cell Biol. 105:3087-3096 (1987)). By this way, syndecan has been proposed to be involved in the maintenance of epithelial morphology. Mouse mammary tumor cells (S115), when steroid-induced to modulate their morphology from an epithelial to a more fibroblastic or fusiform phenotype, lose syndecan expression (Leppä et al., Cell Regul. 2:1-11 (1991)), as to several other cell types in vivo when they transform (Inki et al., Am. J. Pathol. 139:1333-1340 (1991); Inki et al., Lab. Invest. 66:314-323 (1992)), suggesting a general loss of syndecan expression during malignant transformation.

SUMMARY OF THE INVENTION

It is therefore an object of the subject invention to provide a method of altering the differentiated state of a host cell by altering syndecan expression in such cell.

It is a further object of the invention to provide a method to induce and regulate syndecan expression, especially in cells which exhibit a malignant phenotype, regardless of the origin of transformation.

It is a further object of the invention to provide a treatment for the reduction of tumor growth in a patient in need of such treatment, by administration of a composition to such patient, such composition comprising efficacious amounts of one or more agents that stimulate syndecan synthesis in the tumor cells of such patient.

It is a further object of the invention to provide the DNA sequence and localization of promoter, suppressor and enhancer elements for the syndecan gene.

It is a further object of the invention to provide a method for the enhancement of syndecan expression in a host cell, by enhancing syndecan gene transcription.

It is a further object of the invention to provide a method for the enhancement of syndecan expression in malignant cells, by preventing suppression of syndecan gene transcription.

It is a further object of the invention to provide a biochemical method for the inactivation of suppressors of syndecan gene expression in malignant cells.

It is a further object of the invention to provide a method for the stimulation of cellular differentiation by enhancing syndecan expression in both malignant and normal cells.

It is a further object of the invention to provide a method for the stimulation of cellular proliferation and differentiation, thus tissue regeneration, especially in processes such as wound healing, by enhancing syndecan expression.

Further features, objects and advantages of the present invention will become more fully apparent from a detailed consideration of the following description of the subject invention when taken together with the accompanying drawings.

BRIEF DESCRIPTION OF THE DRAWINGS

- Figure 1. Figure 1 is a diagram of the assembly of mouse syndecan gene and its promoter region.
- Figure 2. Figure 2 is the complete sequence of the mouse syndecan gene [SEQ ID Nos. :1: (DNA) and :2: (protein)] and its proximal promoter. Regulatory sites for the expression of syndecan may also exist on the first intron following the first exon (see Figure 1).
- Figure 3. Figure 3 is a diagram of the assembly of mouse syndecan promoter region and the localization of the enhancer and suppressor elements together with restrictions sites for three different enzymes.
- Figure 4. Figure 4 is the complete sequence of the mouse syndecan enhancer element [SEQ ID No. :3: (DNA)].located 8-10 kbs upstream from the transcription initiation site as indicated in figure 3.
- Figure 5. Figure 5A (panels A-D) is a photographic presentation of reduced growth ability of syndecan-transfected cells in soft agar. Panel A (wild-

type S115 cells) and B (control transfected cells) are pictures of the colonies that are formed in soft agar in the presence of testosterone, a feature typical for hormone-transformed cells. This growth ability was not observed with two independent syndecan-transfected cell clones (panels C and D), demonstrating how syndecan re-expression can overcome the effect of hormone-induced transformation. Figure 5B is a graphical presentation of how syndecan-transfected cells lose their ability to form tumors in nude mice. Wild-type or control transfected cells produce tumors in testosterone-administered nude mice while syndecan transfected cells revealed a very low tendency to produce tumors.

Figure 6 is a graphical representation of enhanced syndecan expression in 3T3 cells by simultaneously administered basic fibroblast growth factor (bFGF) and transforming growth factor beta (TGF-β). This is an example of how syndecan expression can be enhanced as a result of growth factor action in normal cells during the differentiation process.

Figure 7 is a graphical representation of enhanced syndecan expression by MCF-7 cells exposed to the anti-estrogen toremifene. When exposed to estrogen, syndecan expression in MCF-7 cells was reduced and the cells transformed. Subsequent treatment with the anti-estrogen (toremifene) restored syndecan expression to levels close to that found in cells not exposed to estrogen and aided the cells in maintaining their normal growth behavior.

Figure 8 is a graphical presentation of how the suppressor element (see figure 3) is active in S115 cells treated with testosterone. Indicated stretches of promoter sequences were transfected in hormone-treated S115 cells and analysed for their transcription activity as described in Example VI. A dramatic drop was observed with suppressor construct as indicated in Figure 3, and it was more obvious in transformed S115 (a) cells than in 3T3 cells (b).

Figure 9 is a graphical presentation of how the enhancer element is active in growth hormone-treated 3T3 cells. Various stretches of promoter were transfected in 3T3 cells and analysed for their transcription activities. Fragment pXb6, which is the same as illustrated in Figure 3 as an enhancer, revealed more than ten fold stimulation in 3T3 cells exposed to growth factors bFGF and TGFβ if compared to non-treated cells.

DEFINITIONS

In order to provide a clearer and more consistent understanding of the specification and claims, including the scope to be given such terms, the following definitions are provided.

"Enhancement" or "Stimulation" of Syndecan expression. By "enhancement" or "stimulation" of syndecan expression" is meant an effect of increasing the synthesis of syndecan, either by the induction or de-suppression (de-repression) of syndecan gene transcription and/or translation.

<u>Cell growth</u>. By "cell growth" is meant cell replication, both controlled and uncontrolled.

Malignant. By "malignant" is meant uncontrolled cell growth.

More Differentiated Phenotype. In stating that a cell has a "more differentiated phenotype" is meant that the cell possesses a phenotype usually possessed by a certain cell type more differentiated than the cell, which the cell was deficient in prior to enhancement of syndecan expression according to the invention. This phenotype may be defined by one or more phenotypic characteristics. For example, an epithelial cell is a more differentiated phenotype of a mesenchymal-like shape; therefore, the ability of the method of the invention to maintain cells in an epithelial cell morphology rather than a mesenchymal-like shape is a more differentiated phenotype within the meaning of the definition. Continuous syndecan expression is necessary for the maintenance of terminal differentiation of epithelial cells.

Syndecan expression is also linked to the normal differentiation of mesenchymal cells. However, only a transient increase in syndecan expression is needed for normal differentiation in mesenchymal cells. Therefore, contrary to epithelial cells, expression of syndecan is not needed for maintenance of terminal differentiation in mesenchymal cell. To induce differentiation of a suitable mesenchymal precursor cell population (such as a "condensing mesenchymal" cell population) to a fully differentiated mesenchymal cell, there is needed only a transient expression of syndecan expression. Therefore, a terminally differentiated mesenchymal cell is a "more differentiated phenotype" than a condensing mesenchymal cell.

Other useful phenotypes that are present in syndecan-deficient cells and not in cells expressing sufficient syndecan include fusiform shapes with long filopodial extensions with extensive under-and overlapping of these processes (so that the cells appear to have a defect in cell adhesion).

in another example, syndecan-deficient NMuMG cells continue to secrete milk fat globule antigen (and thus appear mammary-like) and continue to express cytokeratins (thus appear epithelial-like). However, their actincontaining cytoskeleton is disorganized and their expression of beta1 integrins and E-cadherins at the cell surface is markedly reduced. Upon expression of sufficient syndecan, these phenotypes are corrected so that there is no reduction in cell surface integrins or E-cadherin and the cell has an epithelial morphology. Therefore, the amount of cell surface integrins or E-cadherin are useful markers of syndecan expression and may be used to monitor what amount of a drug is needed for efficacious results according to the method of the invention.

Efficacious Amount. An "efficacious amount" of an agent is an amount of such agent that is sufficient to bring about a desired result, especially upon administration to an animal or human.

Administration. The term "administration" is meant to include introduction of agents that induce syndecan expression into an animal or human by any appropriate means known to the medical art, including, but not limited to, injection, oral, enteral and parenteral (e.g., intravenous) administration.

Pharmaceutically Acceptable Salt. The term "pharmaceutically acceptable salt" is intended to include salts of the syndecan-inducing agents of the Invention. Such salts can be formed from pharmaceutically acceptable acids or bases, such as, for example, acids such as sulfuric, hydrochloric, nitric, phosphoric, etc., or bases such as alkali or alkaline earth metal hydroxides, ammonium hydroxides, alkyl ammonium hydroxides, etc.

Pharmaceutically Acceptable Vehicle. The term "pharmaceutically acceptable vehicle" is intended to include solvents, carriers, diluents, and the like, which are utilized as additives to preparations of the syndecan-inducing agents of the invention so as to provide a carrier or adjuvant for the administration of such compounds.

<u>Treatment.</u> The term "treatment" or "treating" is intended to include the administration of compositions comprising efficacious amounts of syndecan-

inducing agents to a subject for purposes which may include prophylaxis, amelioration, prevention or cure of a medical disorder, including tumor growth, hair growth.

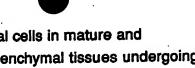
DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENTS

Briefly, in its broader aspects the present invention comprehends a method for maintaining a differentiated phenotype in a normal (non-malignant) cell that otherwise would suppress syndecan expression, by maintaining syndecan expression in such cell. The invention also comprehends a method for inducing a more differentiated phenotype in a malignant cell that lacks (or is deficient in) syndecan expression, by stimulating syndecan expression in such a cell. As used herein, if a cell is said to "lack" syndecan expression, it is meant to include cells that are either completely deficient in syndecan protein or produce insufficient syndecan levels to maintain or attain a desired differentiated phenotype.

The methods of the invention will not only prevent the progression (worsening) of a transformation state and tumor growth of cells, but can also be used to maintain differentiated cells in their differentiated state so that they continue to perform differentiated functions characteristic of such cells. Examples of differentiated functions of non-malignant cells include the secretion characteristics of a cell (that is, the secretion of specific proteins and/or other macromolecules) and hair formation by epidermal cells of skin. Thus, according to the invention, administration of agents capable of inducing syndecan expression in epidermal skin cells of the scalp will promote hair growth among bald (or balding) people.

The subject method may be accomplished in a variety of manners including biochemical, chemical or even molecular biological type methods. While the method is applicable to a variety of cancer (both malignant and non-malignant) and normal cells, it is particularly adaptable for treating malignant cells which have become transformed. This includes cells transformed due to hormonal influences of the body or environmental influences, such as chemicals or radiation exposure. Especially, the tumor type is a tumor characterized by loss of syndecan, for example, a glioma, myeloma, carcinoma, sarcoma, lymphoma, or adenoma.

Generally, any cell genetically capable of expressing syndecan can be a cell stimulated to express syndecan by the method of the invention. Syndecan-



is naturally expressed in a wide variety of epithelial cells in mature and embryonic tissues and by various embryonic mesenchymal tissues undergoing inductive interactions with epithelia. In addition, syndecan is naturally expressed on Leydig cells and on developing B-lymphocytes and a subpopulation of plasma cells.

Enhanced syndecan expression may be achieved by administration of compositions containing a biochemically, and/or chemically and/or molecularbiologically active component to an individual. Compositions may be administered orally, intravenously, subcutaneously or locally, or by any other method which will allow cells, normal or malignant, to be exposed to syndecan expression enhancing component.

By a "biochemically" or "chemically" active component is meant a component that alters the endogenous syndecan biochemistry or chemistry of the target cell without altering syndecan gene expression per se. For example, such alterations may include altering the half-life of syndecan protein or mRNA, so as to increase levels of syndecan protein in the cell, for example, by altering the external domain of the cell's endogenous syndecan, or the cell surface membrane properties in general, so as to retain higher levels of syndecan on the cell surface; and, altering the syndecan protein active site(s), so as to enhance the efficiency of the syndecan response.

By a "molecular-biologically" active component is meant a component that alters endogenous syndecan gene expression in a manner that allows for an increase in cellular syndecan, such as, for example, by stimulating transcription, preventing (or reducing) suppression of transcription, derepression transcription, and generally increasing levels of mRNA and/or translation efficiency.

It is known that cellular transformation involves activation of some cell growth stimulating genes (like oncogenes) and inactivation of some other genes, which work to suppress cell growth. It has recently been shown that loss of syndecan expression is observed upon transformation of cells, and that this suppression is due to syndecan gene inactivation (Leppä et al., Cell Regul. 2:1-11 (1991); Inki et al., Am. J. Pathol. 139:1333-1340 (1991); Inki et al., Lab. Invest. 66:314-323 (1992)). This was demonstrated in several biological models of various known carcinogenic systems (oncogenes, chemical carcinogens, UV-light, hormone-exposure, etc.). Therefore, syndecan gene suppression is implicated in the development of cellular transformation.

Further, according to the invention, all the manipulations of such cells which can induce syndecan expression in malignant cells, also induce these cells to obtain a more differentiated phenotype, and thus, subsequently reduce their potential tumorigenic behavior and metastasis.

In a preferred embodiment, the cell in which syndecan expression is stimulated is a cell that is steroid-responsive. Examples of such steroid-responsive cells include breast cells, endometrium cells and prostate cells, especially in the malignant state. In a highly preferred embodiment, the cell is responsive to estrogen and/or androgen.

Examples of other cell types that will respond to the treatment of the invention include malignant and non-malignant mesenchymal cells.

The regulatory elements of a given gene are commonly located upstream from (5' to) the transcription initiation site. Syndecan, however, has a very peculiar gene structure, in which the first and second exons are separated by a very large intron (Figure 1). This could mean that, in addition to the base sequences upstream from the transcription site, syndecan expression may also be susceptible to regulation by base sequences located between first and second exons (Figure 2), that is, in the first intron.

It has now surprisingly been found that the syndecan gene has a strong enhancer element located 8-10 kb upstream from the transcription initiation site. The sequence of this enhancer element has been identified and is given in Figure 4 [SEQ ID. No. 3].

Manipulation of the upstream region of the syndecan gene can block its inactivation during malignant transformation. For example, replacement of the region in front of first exon of the syndecan gene with the glucocorticoid-inducible elements of mouse mammary tumor virus (MMTV) not only blocks syndecan suppression during malignant transformation, but also Inhibits the ability or potential of cells to transform and become tumorigenic (Figure 3). These findings suggest a very important role for syndecan in the maintenance of normal epithelial morphology (Leppä et al., Proc. Natl. Acad. Sci. USA 89:932-936 (1992)).

Cells destined to differentiate during organ formation or tissue regeneration also exhibit enhanced syndecan expression (Vainio *et al., Dev. Biol. 147*:322-333 (1991); Elenius *et al., J. Cell Biol. 114*:585-595 (1991)). The component(s) responsible for the regulation of syndecan expression (either

directly or indirectly) have not yet been identified. Growth factors are candidates for this role since they are known to be involved in the regulation of early development and cellular differentiation (Heath *et al.*, *Curr. Opin. Cell Biol. 3*:935-938 (1991)). Involvement of growth factors is also supported indirectly by the fact that the expression of two embryonally important growth factors (TGF-B and FGF) has been shown to coincide with syndecan within developing tooth (Vaahtokari *et al.*, *Development 113*:985-994 (1991); Wilkinson *et al.*, *Development 105*:131-136 (1989)).

Based on these findings the possible effect of growth factors on the expression of syndecan has been tested. It was shown that both bFGF and TGF-B alone, but especially if administered in combination, enhance syndecan expression by 3T3 cells (Figure 4). This stimulation is close to the levels observed in syndecan-expressing epithelial cells (Elenius *et al.*, *J. Biol. Chem. 267*:6435-6441 (1992)) prior to their malignization (Leppä *et al.*, *Proc. Natl. Acad. Sci. USA 89*:932-936 (1992)). These findings suggest that growth factors, and their derived fragments and domains may prove to be valuable for the development of an active tool for the regulation of syndecan expression.

Preferably, for treatment of humans and animals, a drug is administered that results in the enhancement of syndecan expression to levels sufficient to facilitate cellular differentiation in the degenerative stages of tissues. Such drugs are herein termed "syndecan-inducing agents." Syndecan-inducing agents include the growth factors and their derivatives that retain growth-factor activity. Examples of such growth factors include bFGF, and TGF-B, whether administered separately or together.

Even more preferred is a syndecan-inducing agent that has good tissue and cell penetration so that it could directly interfere with suppressor(s) of syndecan expression within cell nuclei. Such a syndecan-inducing agent is the known antitumor drug toremifene. When toremifene, known to have good plasma membrane penetration, is administered to the hormone-transformed epithelial cells with reduced syndecan expression, the cells reverse their lowered syndecan expression, and induce syndecan levels to those close to the level observed with normal, non-transformed cells (Figure 5). This demonstrates that syndecan-inducing agents useful in the methods of the invention are known and available and that such agents can specifically prevent cellular malignization by blocking suppression of syndecan expression. Another useful drug in this regard is tamoxifen.

Such syndecan-inducing agents may be administered using currently available preparations, or in any pharmaceutically acceptable vehicle. The route of administration may be any route that delivers efficacious levels of the drug to the desired active site, for example, by injection.

For parenteral administration, preparations containing one or more syndecan-inducing agents may be provided to the patient in need of such treatment in combination with pharmaceutically acceptable sterile aqueous or non-aqueous solvents, suspensions and emulsions. Examples of non-aqueous solvents are propylene glycol, polyethylene glycol, vegetable oil, fish oil, and injectable organic esters. Aqueous carriers include water, water-alcohol solutions, emulsions or suspensions, including saline and buffered medical parenteral vehicles including sodium chloride solution, Ringer's dextrose solution, dextrose plus sodium chloride solution, Ringer's solution containing lactose, or fixed oils. Intravenous vehicles include fluid and nutrient replenishers, electrolyte replenishers, such as those based upon Ringer's dextrose and the like.

The syndecan-inducing agent of the invention can be employed in dosage forms such as tablets, capsules, powder packets, or liquid solutions for oral administration if the biological activity of the syndecan-inducing agent is not destroyed by the digestive process and if the characteristics of the compound allow it to be absorbed across the intestinal tissue.

The syndecan-inducing agents may also be administered by means of pumps, or in sustained-release form. The syndecan-inducing agents used in the method of invention may also be delivered to specific organs in high concentration by means of suitably inserted catheters, or by providing such molecules as a part of a chimeric molecule (or complex) which is designed to target specific organs.

Administration in a sustained-release form is more convenient for the patient when repeated injections for prolonged periods of time are indicated.

The composition containing the syndecan-inducing agent can be manufactured in a manner which is in itself know, for example, by means of conventional mixing, granulating, dragee-making, dissolving, lyophilizing or similar processes. The compositions of the present invention that provide the syndecan-inducing agent, in and of themselves, find utility in their ability to slow or prevent tumor growth or tumor reappearance. The syndecan-inducing

compositions of the invention utilize the body's own mechanisms for promoting differentiation of specific cell types to its maximum potential.

In intravenous dosage form, the compositions of the present invention have a sufficiently rapid onset of action to be useful in the acute management of tumor growth.

Additionally, a low potency version is useful in the management of disorders wherein a tumor has been effectively treated and the patient appears to be in remission, but it is desired to maintain sufficient levels of syndecaninducing agents in the patient so as to assist the body in preventing a recurrence of the tumor.

Typical doses of toremifene or tamoxifen, and other such syndecaninducing agents useful in the methods of the invention for treatment of humans or other animals are 20-600 mg daily, and preferably 20-60 mg daily.

The examples below are for illustrative purposes only and are not deemed to limit the scope of the invention.

EXAMPLE 1

Reversal of hormone-induced transformation by exogenous syndecan expression.

As previously described (Leppä et al., Cell Regulation 2:1-11 (1991)), S115 mouse mammary tumor cells were routinely cultured in DMEM. For experimental studies involving hormone treatment, inactivated fetal calf serum (i-FCS) was replaced with 4% dextran charcoal-treated fetal calf serum (DCC-FCS), which eliminates endogenous steroids from serum), and used either with or without testosterone (10 nM) and with or without dexamethasone (10 nM or 1 μM). Cells were plated at a density of 10,000 cells/cm 2 and the medium was replenished every 3 days.

Plasmid pUC19-hsynpr7 containing human syndecan cDNA (Mali et al., J. Biol. Chem. 265:6884-6889 (1990)) was digested with Nael restriction endonuclease, and the derived 336 bp long-fragment was separated in and eluted from low melting agarose gel. Plasmid pUC19-hsyn4 (Mali et al., J. Biol. Chem. 265:6884-6889 (1990)) was digested with Nael and Hindll (polylinker site), and the plasmid-containing fragment starting from base 487 was isolated. The Nael fragment from hsynpr7 was ligated to the pUC-hsyn Nael/Hindll

digested vector. The orientation of insert was verified by restriction enzyme analysis and sequencing. The derived plasmid, containing the full coding region of human syndecan core protein, was named pUC19-hsynfull. This plasmid was further digested with *Bam*HI and *Sph*I (polylinker site). A fragment containing syndecan coding region bases 150-1461 was isolated and bluntended, using Klenow and T4 DNA polymerase. Finally, this fragment was ligated to *Sal*I-linearized and bluntended pMAMneo vector (Clontech; Palo Alto, CA), resulting in a chimeric gene containing RSV-MMTV-LTR promoter connected to human syndecan coding region and SV-40 polyadenylation signal. The orientation was confirmed by restriction enzyme digestions. The plasmid was named pMAMneo-hsyn.

For control transfections, a 642 bp long-*Hind*III/*Pvu*II fragment of human growth hormone gene (consisting of exons 4 and 5; Bornstein *et al., J. Biol. Chem. 263*:1603-1606 (1988)) was blunt-ended and cloned into the same pMAMneo vector, as described above. This control construct was named pMAMneo-hGH.

All plasmids were isolated using the CsCl density gradient method. Before transfections, plasmids were linearized with *Mlu*l, chloroform/phenol extracted and ethanol precipitated.

Transfections were performed using Lipofectin[™] (BRL), according to manufacturer's Instructions. After selection for two weeks (G418; 750 μg/ml, Sigma), surviving clones were isolated from growth plates using cloning cylinders. The expression of human syndecan or growth hormone (consisting of exons 4 and 5) mRNAs was then confirmed by RNA isolation and Northern blot analysis. Clones expressing high levels of transfected genes were selected for further studies and characterizations. These stock cells were routinely cultured in the presence of G418 (300 μg/ml).

For the measurement of exogenous syndecan expression total RNA was isolated from wild-type S115 cells and cells transfected with human syndecan or growth hormone genes. RNA was extracted using 4M guanidine isothiocyanate and CsCl pelleting, as earlier described by Chirgwin *et al.*, *Biochemistry 18:*5294-5299 (1979)). RNA from normal mouse mammary NMuMG and normal human mammary HBL-100 cells was used for comparison. RNA aliquots of 15 μg were separated in 1% formaldehyde agarose-gel electrophoresis and transferred to GeneScreen PlusTM hybridization membrane (New England Nuclear). Blots were hybridized with multiprime (Amersham)

labeled inserts of either mouse (PM-4) (Saunders et al., J. Cell Biol. 108:1547-1556 (1989)) or human syndecan (pUC19-hsyn4 BamHI 1.1 kb fragment) (Mali et al., J. Biol. Chem. 265:6884-6889 (1990)), or with human growth hormone exons 4 and 5 (hGH) (Leppä et al., Proc. Natl. Acad. Sci. USA 89:932-936 (1992)) cDNAs, using the high stringency conditions suggested by the manufacturer of the membrane (New England Nuclear). All techniques based on modern molecular biology are fully explained in the literature such as in the laboratory manual entitled Current Protocols in Molecular Biology.

Anchorage independent cell growth was measured in a soft agar colony assay. The six well-plates were first covered with an agar layer consisting of 2 ml DMEM, 0.5% agar and 4% DCC-FCS. The middle layer contained 10⁴ cells in 0.5 ml DMEM supplemented with 0.33% agar and 4% DCC-FCS, with or without 10 nM testosterone. The uppermost layer, consisting of medium (2 ml), was added to prevent drying of the agarose gels. The plates were incubated at 37°C in 5% CO₂ for 12 days after which cultures were evaluated and photographed.

Tumorigenicity of S115 wild type cells, one hGH transfected control clone and two clones expressing human syndecan-1 was measured in nude mice. For this cells were cultured in DMEM containing 5% FCS and 10 nM testosterone. After four days in culture, cells were harvested with trypsin, washed, and 10⁷ cells suspended in 0.2 ml of DMEM were injected subcutaneously into back of athymic male nude mice (balb-C). Silastic testosterone capsule, which is known to increase the growth rate of S115 cells (King et al., J. Steroid. Biochem. 7:869-873 (1976)) was simultaneously implanted. Nude mice were examined regularly for tumor development and the size of the palpable tumors measured at intervals.

EXAMPLE II

Growth factors enhance syndecan expression.

NMuMG mouse mammary epithelial cells and 3T3 (NIH) mouse fibroblasts were routinely cultured in bicarbonate-buffered Dulbecco's modified Eagle's medium (DMEM; GIBCO) containing 10% FCS (GIBCO) and antibiotics, as previously described (Elenius *et al.*, *J. Biol. Chem. 265*:17837-17843 (1990)). For experiments, cells were plated at equal density on culture dishes (Nunc) and grown to 60 - 70% confluency. Twenty-four hours before supplementing growth factor(s) to the medium, fresh medium containing 2%

CMS-FCS (Vogel *et al.*, *Proc. Natl. Acad. Sci. USA 75*:2810-2814 (1978)) was replaced. Equally treated cultures without growth factor addition served as negative controls. Porcine TGF61 (R&D), recombinant human bFGF (Boehringer) and murine EGF (Sigma) were used in final concentrations of 2.5 ng/ml (100 pM), 10 ng/ml (570 pM) and 1.2 ng/ml (200 pM) respectively, in all experiments. For quantitation and isolation of cell surface syndecan, media were discarded at time points indicated in the text and the cell layers were washed twice with ice cold phosphate buffered saline (PBS). Cells were scraped with a rubber policeman into ice cold PBS supplemented with 0.5 mM EDTA and centrifuged. After subsequent washes by resuspension and centrifugation the cell numbers were measured by counting the nuclei with a Coulter Counter (Coulter Electronics).

For quantitation of syndecan Intercalated into the cell membrane, syndecan ectodomain was released by incubating washed cells in 20 μg/ml bovine pancreatic trypsin (Type III; Sigma) in PBS for 10 min on ice bath. After incubation the cells were centrifuged, leaving the ectodomain in the supernatant (Rapraeger et al., J. Biol. Chem. 260:11046-11052 (1985)). Sample volumes equal to 400,000 or 200,000 cells for 3T3 or NMuMG cells, respectively, were loaded on cationic nylon membrane (Zeta-Probe; BioRad) in a minifold-slot apparatus (Schleicher and Schuell), as previously described (Jalkanen et al., J. Cell Biol. 105:3087-3096 (1987)). Nonspecific binding was blocked by incubating the membrane for one hour at room temperature in PBS containing 10% FCS. Syndecan attached to the membrane was detected with a monoclonal antibody against mouse syndecan core protein (mAB 281-2) (Jalkanen et al., J. Cell Biol. 101:976-984 (1985)) that was radiolodinated by chloramine-T oxidation method (Stähli et al., Meth. Enzymol. 92:242-253 (1983)). The membrane was incubated overnight at 4°C with 1251-labeled 281-2 in PBS + 10% FCS (10,000 CPM/ml). After five washes with PBS the bound antibody was visualized by autoradiography.

The accumulation of syndecan ectodomain into the medium was estimated by taking samples corresponding to 1/50 (3T3 cells) or 1/100 (NMuMG cells) of the total volume of the remaining medium at selected time points. The samples were analyzed by loading them to nylon membrane as described above. The autoradiography signal was quantitated with a GelScan XL ultroscan densitometer (LKB) using GelScan XL 2400 software (LKB).

For syndecan purification, cells were radiolabeled for 24 hours in low sulfate DMEM (MgCl $_2$ substituted for MgSO $_4$; 2% CMS-FCS) with 100 μ Ci/ml

35_{SO4} (New England Nuclear) in the presence or absence of growth factor(s). Cell surface trypsin-releasable material was collected, as described above, and after dialysis against Tris-buffered saline (TBS), the sample was loaded onto a 281-2-Sepharose CL-4B immunoaffinity column (Jalkanen *et al.*, *J. Cell Biol.* 105:3087-3096 (1987)). Bound material was eluted with 50 mM triethylamine (TEA) (pH 11.5) and the amount of radioactive PG in each fraction was analyzed using cetylpyridiumchloride-impregnated Whatman 3MM filter discs (Rapraeger *et al.*, *J. Biol. Chem. 260*:11046-11052 (1985)). For interaction experiments, fractions containing most of the labeled PG were pooled and dialyzed against PBS.

To obtain unlabeled syndecan ectodomain for interaction assays (see below) the same procedure was used except that no radioactive sulfate was added to the culture medium and the syndecan containing fractions eluted from the immunoaffinity column were detected by immuno-dot assay using mAB 281-2. The estimation of the molar concentration of syndecan was based on the use of previously determined syndecan concentration by total amino acid analysis (Jalkanen et al., J. Cell Biol. 106:953-962 (1988)).

SDS-PAGE and Western Blot - For western blot experiments cells were cultured 24 hours with or without growth factor(s). Syndecan ectodomain containing material released from the cell surface by trypsin treatment was fractionated on SDS-PAGE gradient (2 -15%) gel (O'Farrel, J. Biol. Chem. 250:4007-4021 (1975)). After electrophoresis, samples were transferred onto Zeta-Probe membrane using electroblotting 2005 Transphor apparatus (LKB). The syndecan antigen on the filter was detected with radiolodinated mAB 281-2 and the filter was washed, as described above for slot blot analysis.

Northern Blot - RNA was isolated from 3T3 and NMuMG cells using 4 M guanidine isothiocyanate and CsCl density centrifugation (Chirgwin et al., Biochemistry 18:5294-5299 (1979)). RNA samples were size-separated on 1% agarose formaldehyde gel, transferred to GeneScreen Plus™ membrane (New England Nuclear) and hybridized with multi-prime (Amersham) labeled partial cDNA clone for mouse syndecan (PM-4) (Saunders et al., J. Cell Biol. 108:1547-1556 (1989)). After hybridization the membrane was washed in 2 x SSC and 1.0% SDS at 65°C (high stringency conditions). For rehybridization with glyceraldehyde-3-phosphate-dehydrogenase (GAPDH; Fort et al., Nucleic Acid Res. 13:1431-1442 (1985)) the bound PM-4 probe was removed as recommended by the manufacturer of the filter (NEN).

EXAMPLE III

Induction of syndecan mRNA expression in the human breast cancer cells (MCF-7) growth-inhibited with toremitene.

The steroid-responsive human breast cancer cell line MCF-7 was used to study the expression of human syndecan under different growth conditions regulated by estrogen and antiestrogen. For the experiment cells were plated at a density of 1.2 x 10⁶ cells/100 mm ø plastic culture dishes and grown as monolayer cultures in 10 ml per dish of phenol red-free DMEM medium with 5% dextran/charcoal stripped fetal calf serum (DS-FCS), 2 mM L-glutamine and 3 μg/ml insulin. For hormone-treatment 1 nM 17β-O-estradiol (E2), alone or with 1-6.25 μM toremifene, dissolved 70% in ethanol, were added to the culture medium on the day following plating. The cells were cultured for 6 days, and the media were changed every second day. For RNA extraction the cells were washed *in situ* with PBS and scraped from the plates in 4 M guanidine isothiocyanate.

EXAMPLE IV

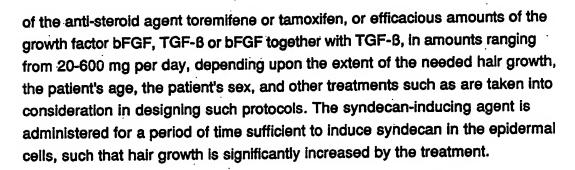
Treatment of Steroid-Responsive Tumors in Patients.

Patients diagnosed as having a steroid-responsive tumor selected from a breast tumor, an endometrium tumor, a prostate gland tumor or a mesenchymal tissue tumor are administered a composition that contains efficacious amounts of the anti-steroid agent toremifene or tamoxifen, or efficacious amounts of the growth factor bFGF, TGF-B or bFGF together with TGF-B, in amounts ranging from 20-600 mg per day, depending upon the extent of the tumor, the patient's age, the patient's sex, and other treatments such as are taken into consideration in designing such chemotherapeutic protocols. The syndecan-inducing agent is administered for a period of time sufficient to induce syndecan in the tumor cells, such that the tumor cells now take upon a more differentiated phenotype and such that the growth of the tumor is arrested or significantly slowed by the treatment.

EXAMPLE V

Stimulation of Hair Growth in Epidermal Skin Cells.

Patients diagnosed as being in need of increased hair growth in the scalp region are administered a composition that contains efficacious amounts



EXAMPLE VI

Dermination of Mouse Syndecan Promoter and Enhancer Activities

The mouse syndecan gene has been cloned and characterized up to -10 kbs upstream from the transcription start site. To determine the specific activities of different proximal promoter regions (up to -2 kbs from the start site) and enhancer regions (from -2 to -10 kbs) we have made plasmid constructs where these regions were cloned into pCAT basic or pCAT promoter vectors, containing the CAT reporter gene. The reporter CAT gene produces chloramphenicol acetyltransferase enzyme, which transfers the n-butyryl moiety of n-butyryl CoA to chloramphenicol. The n-butyryl chloramphenicol can be separated from native chloramphenicol by xylene.

For the further characterization of the proximal syndecan promoter a series of retriction enzyme treatments was made on the upstream region (Hind III, Hind II, Bgl II, Stu I, Dra I, Cla I, BamHI and Pst I — Xho I) and the obtained fragments were cloned into the pCAT basic vector. For enhancer areas, three Xba I fragments were cloned into a pCAT promoter vector, where the SV 40 promoter was displaced by the Bgl II - Xho I fragment from the syndecan promoter.

The plasmid constructs were transiently transfected into eukaryotic cells by calcium phosphate precipitation simultaneously with a β-Galactosidase expressing vector to determine the transfection efficiency. After a four hour incubation cells were treated with 15% glycerol and grown for approximately 48 h in cell culture medium. Cells were then scraped from dishes in TEN-buffer and the cytoplasmic extract was obtained by repeated freezing and thawing. β-Galactosidase acitivity was obtained in hte cytoplasmic extreact by adding 0.1 M sodium phosphate, 45 mM mercaptoethanol and 0.2 mg ONPG. This was incubated from 2 hours to overnight and the colour reaction was measured spectrofotometrically at 420 nm.

The CAT activity was determined by adding 0.25 M Tris buffer, 25 ng n-butyryl CoA and 0.0626 μ Ci of 14 C-chloramphenicol to the cytoplasmic extract. These were incubated overnight, backextracted with xylene and radioactivity was measured by scintillation counting. The CAT activity was corrected for transfection efficiency by β -galactosidase activity.

The cells used for proximal promoter constructs were 3T3 NIH, S115 (either hormone-treated of not) and nMuMG cells. For enhancer constructs we used 3T3 NIH cells grown in 2% CMS medium and to test the effect of growth factors in 2% CMS medium with 10 ng/ml FGF-2 and 2ng/ml TGFB-1.

All references cited herein are fully incorporated herein by reference. Having now fully described the invention, it will be understood by those with skill in the art that the scope may be performed within a wide and equivalent range of conditions, parameters and the like, without affecting the spirit or scope of the invention or any embodiment thereof.

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT:
 - (A) NAME: Alanen-Kurki, Leena
 - (B) STREET: Piispankatu 6 E 31
 - (C) CITY: TURKU
 - (E) COUNTRY: FINLAND
 - (F) POSTAL CODE (ZIP): FIN-20500
 - (A) NAME: Auvinen, Petri
 - (B) STREET: Kirschgartenstrasse 5
 - (C) CITY: HEIDELBERG
 - (E) COUNTRY: GERMANY
 - (F) POSTAL CODE (ZIP): D-69126
 - (A) NAME: Jaakkola, Panu
 - (B) STREET: Kellonsoittajankatu 13 B 20
 - (C) CITY: TURKU
 - (E) COUNTRY: FINLAND
 - (F) POSTAL CODE (ZIP): FIN-20500
 - (A) NAME: Jalkanen, Markku
 - (B) STREET: Rauvolantie
 - (C) CITY: PIISPANRISTI
 - (E) COUNTRY: FINLAND
 - (F) POSTAL CODE (ZIP): FIN-20760
 - (A) NAME: Leppa, Sirpa
 - (B) STREET: Valkintie 14 as 6
 - (C) CITY: LITTOINEN
 - (E) COUNTRY: FINLAND
 - (F) POSTAL CODE (ZIP): FIN-20660
 - (A) NAME: Mali, Markku
 - (B) STREET: Inkereentie 176
 - (C) CITY: SALO
 - (E) COUNTRY: FINLAND
 - (F) POSTAL CODE (ZIP): FIN-24280
 - (A) NAME: Vihinen, Tapani
 - (B) STREET: Kaskenkatu 11 C 54
 - (C) CITY: TURKU
 - (E) COUNTRY: FINLAND
 - (F) POSTAL CODE (ZIP): FIN-20700
 - (A) NAME: Warri, Anni
 - (B) STREET: Juhaninkuja 2 as 2
 - (C) CITY: LIETO
 - (E) COUNTRY: FINLAND
 - (F) POSTAL CODE (ZIP): FIN-21420
- (ii) TITLE OF INVENTION: SYNDECAN STIMULATION OF CELLULAR DIFFERENTIATION
- (iii) NUMBER OF SEQUENCES: 3



- (A) MEDIUM TYPE: Floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
- (v) CURRENT APPLICATION DATA:

APPLICATION NUMBER: WO TO BE ASSIGNED

- (vi) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 07/988427
 - (B) FILING DATE: 01-DEC-1992

(2) INFORMATION FOR SEQ ID NO: 1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26700 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: both
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iii) ANTI-SENSE: NO

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: join(4378..4443, 22026..22106, 23001..23483, 23905..24039, 24251..24418)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

60	GAGTGACTAG	CTATTGGTGG	ATGTCCACCC	AGATGGAGTG	CAAACTCACC	TCTAGATATT
120	CCCTCACACC	TTAGGACCCA	AGCTAGCTCT	TCAGATGCTT	GTCTTCTGAC	TCTTTCCTCT
180	GACATGAGAA	CAGTGGAGTT	ACCTTTAACC	CTCTCTTAGT	ACTTTATTTG	TGCAAATAAT
240	AAATTAAAA	AAAGTAAAAŢ	TCATAAATGA	ATATTTCATT	ATAATTTATA	ATTAACTACC
300	CGCAAGTCCT	GTGGCTCCAA	GTAAAGGCCA	TGGCCCAGTG	TTGAGCATGA	ATAGAAAGGT
360	GAGGCTAAAA	TTTATTGATT	TTGAGGGAAG	TTCTTCAGGC	AACGGGCCTG	GACAAATGGT
420	CACACTGCAT	TGTGCTCTCC	AAGCCCTGGA	GCCTAGTGTG	GGCTCCACTT	GCAACCCAAA
480	AGGTCATTAG	GGGGAGTCCA	TGAGGATGAT	CCTGGGAAGC	GGTGTCAGCA	GTCCACCTGT
540	AATAAGCACA	AAGAAAAAA	GGTCACAAAA	GGGGTACATG	TAGGCTAGCT	CTACATAGTA
600	TAAGACAGCC	GCTGTGAGTT	GGGGGGGATT	AGACCAATGG	AGCACTTGAC	TTGTAATCCC

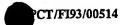
TGGCCTACAA	AGAAAAACCC	TACCCAAACC	CAAGAAAAAT	GAAACCAGTA	ATATAAATAG	660
CTATTTCAT	TTTAAATGCT	CTAAAGACAC	AGCGTTAACA	CAAAAGCTCT	CGTCTGTGGT	720
TCCTATTCCC	TCCTTCTCCC	CCAGGTCTTC	TTTAATGTAT	ACTTTTTGTT	TGCTTATTTG	780
СТТСТТТТСС	ATTTTGGCTT	TTAAAGACAG	GGTCTCACTA	TGTAGCTCCA	ACTATTTGGG	840
AACTCACTAT	GTAGACCAGG	CTAGCCAGGG	ACTTATAGAG	ATCTACCTAC	CACTGCCTCC	900
CAAGTGCTGA	GACTAAAGGC	ATGTGACACT	TTGCTTGGTT	ATTACAAACA	TTTTAAAAGA	960
ACATTTTGAA	CATTAATAGA	TGTATGTATA	TATATCACTC	TATGTAGTAT	ATATGTTAĠA	1020
CATTTTTCAC	TTGAGATACA	TATTTACTCT	CAAAATAAGT	TTTTTGTTTT	TTTTTCTTCT	1080
ТТТДААТТТТ	ATTTTATTTT	TTTTTTTTTT	ATTTTATTAT	TATATGTAAG	TACACTGTAG	1140
CTGTCTTCAG	ACANACCAGA	AGAGGGAGTC	AGATCTTGTT	ACGGATGGTT	GTGAGCACCA	1200
TGTGGTTGCT	GGGATTCGAA	CTCTGGACCT	TCCGAAGAGC	AGTCGGGTGC	TCTTACCCAC	1260
TGAGCCATCT	CACCAGCCCC	TTAAATTTAT	TTTTATCTTA	TGTCCATTGG	TGTTTTGCCT	1320
GCATGTATGT	GTAAAAGTGT	CAGAAACTGA	AGTTACAGAC	TGTTGTGAGC	TACCATTGTT	1380
GTGGGTGCTG	GGACTTGAAC	CTGGGTCCTC	TGGAAGAGCA	GTCATTATTC	TTAACCACTG	1440
AGCCATCTCT	CTAGCCCTCG	TTTTTTAGTT	TTTTTTTTTG	TTTTGTTTTG	TTTTTTGTTT	1500
TTTTAAGATT	TTCTTATTTA	TTATATGTAA	GTACACTGTA	GCTGTCTTCA	GACACTCCAG	1560
AAGAGGGCGC	CAGATCTCGT	TATGGATGGT	TGTGAGCACC	ATGTGGTTGC	TGGGAATTGA	1620
ACTCCAGACC	TTTGGAAGAG	CAGTCAGTGC	TCTTAACTGC	TGAGCCATCT	CTCCAGCCCC	1680
GTTTTTTAGG	TTTTTGAAGA	CAGGGTTTCC	TGTGTAGCTC	TAGCTGTCCA	GGAACTAGCT	1740
CTGTAGACCA	GGTTGGCCTC	AAATTTAGAG	ATTTGCCTGT	CTCTCTGCCT	CTCGAGAGCT	1800
GGGATTAAAA	GTGTGCAGCC	CAACAATCTA	CTCAAAGTAG	GTTTTGAAAA	AGCTTTCCAT	1860
ATTAGGAGTT	AACTAGCTTC	ATTTCAGAAA	TACTGCATGG	AATTCAAATG	TGGGACCATT	1920
CATAGCTACT	TTGGTTTTCC	TTCAGTGACA	GGCATTCGGC	ATGCCTATTA	GGGAAGTCAA	1980
ATGGCCTGGA	GAAGTCATCC	TGGGTGAGAG	GGCTAATGCA	TTTTCAGCTT	GACAGACACT	2040
GTCAACCTAT	GCAGACAGTC	TGCTCCAGCT	CAGATGTCAA	TTGCATGCAG	ACCTGCAGTC	2100
AGACGCTAAG	CTCCCTACCT	ACTCTCCATC	AGCTTAGATG	TAAGGGGTGC	TGGAACAAAG	2160
GCTCTCTCTC	TCTCTCTCTC	TCTCTCTCTC	TCTCTCTCTT	TCTTAGAATT	AGTATTCTAT	2220
TTTATTTTAT	GTAAATTGGT	ACTTCACTTA	CATGTATGTC	CGTGTGAGGA	TGTTGTATCC	2280
TCTGGTACTG	GAGTTATAGA	CAGCTGTAAG	TCGCCATACA	GGTGCTGGGA	ATTGAACCCT	2340

GATCCTCTGG AAGAATAGTC AGTGCTCTTA ACCCCTGAGC CATCTCTCCA ACCTCTTGCA 2400 TATTGAGGAC AGGGAGGAAT CACAAGCCAT GTAGGGTGCC TGGGCTCTGA GGTCAACAGG 2460 ACCATAGCCT CCTTTCTTTA TGTGCCTTTC TTGGGGTCTC CCTATAGGAG TCGTCTTCGT 2520 TGCCTCTTTA CTGTCTCATT GATCTGGGCT AAACTTATGC AGTTGGAAGG AAAGATCAAG 2580 CTGGTCATGT TTAAAACATG AAACAGCCTC ATCAGTTCCC TTCCTGTTCC CGTCTCCCCC 2640 2700 CCCCTCCCG CCCCATTTT GAGAGGACAG GAAGGTAAAA TACCAAAGTG TCCTATTTTC CTCCAAATAT CAGGCTCAAA GGACTGAAGA GCTGACTTCA GATCCCAAAG CCACTGTGTT 2760 AGGAGGCACC TGCTTTTTAG GTCCTAAGCC TTCCTGAGCC TTGCTATTGG GTATTCTTTA 2820 CCAAGACCCT CAAGGATCTA GGCAAGAACT GGGCAGGATC TGTATGTAGC CCATAGTTAG 2880 2940 ACCTAGGGCA GCTGAGACGC CAAAAGGGAG AGTTTCCTGA GGACAAAAGT GTTCAAACAC AACTGGGTGC TGGTTGTTGG GCTACTCGTG GAGGTGTGGT GTGTGTAAAG GAGGCTGTTG 3000 AATTCCCAGA AGGCTGGTTC CACAGTGTAG AGTCTACACT GGGGACTTCC CGAGACGCTG 3060 AGCCTCAGAT CTAGCTTCTC AGTCCAGGCC AGCTGATGTG GGGCTGAGGA ACAAGGATGG 3120 ATGCCATCTA TGGCCCTGCC TTGCAGGTGC AAAGGGCCTT TGGCACCATC TACAGATTGA 3180 GGGCAAGACA GGGCTGGTTC TTCCTCCTTG CTCTCGCTGC TATCTGCCTC GCCTGTAGGC 3240 TCTCTGGGCT CCTTTTTGGA CTGACACGTC TGAAGGAGCT TGGAAACTGT GAGGTCCAGG 3300 CCCCATAGAG AATCATGAAG GAACAGGAAT TCAACTGGAG CTCCGCAGCT GGTTAGGCCT 3360 GCGGTCACCT GGAAACAAG AGGCCATTTA TTTTTTCCTT TGGTCTTGGA CAAGGAAGAG 3420 AAGGGGCTTT CTATAAATAG AAAGACAGCA AAAAAGAAAA TAATAATAAT AATAATAATA 3480 ATAATAATAA TAATAAAAAC AATAACAAAG CCAGCTCTTC CAGACAGTGC TCATGTCTTT 3540 AAAGGTCTTT AAAGGTCTGG AGTTCCCAGC AATTAAGTAA AGGACCAAGA CCTCAGGGGT 3600 CCCCTATCCT CAGCCCGTGG GGAGGTGGGA ACCATACATC GATCCCTCGG TTTATATATA 3660 GCCTCATCGC TGTGGGGCTC CGAGGTTGCC CCCAAAATCT TGCTCACCTG GAGGACCCCT 3720 GGGTGTCCTC GCCCAGAGGG CGCTGCAGCC TCGCACGTAG AGAACTAACA TCGCCCTTCT 3780 CCAGGGCAGT GCCTCCGGAC TCCGGACCAG GACATAGTAG CGAGTGCACC TGGGTCTCCG 3840 TCAGCTACGC ATCAAGGAAG GTGCGACGCG GGAATTACAG ATTGCCGGCA CTCACCAGTG 3900 CTCAGGGGAG GAAGGTGGGA CTCAGACCTG CAAGAGCTGA AGAGTGGGGT GGGCTTCGAT 3960 CCTAGGAGGC GTGGAAGGGG GTGTGGCTGG ATCCCTGGGG GGTGGGGCGC TCCAAGGGGC 4020 GGGGCAACCC AGGGGGCGGG GCCCGAGGGG TGGAGATTGG GACTACCCAG GCCCGCGGAG 4080

•	
CTGGGGGTGG GCGGCTAGTT TTGCAACTGC AGAGCCTTTG GGTTTATTAT AAGGCGGAGC	4140
TCCGCGGGAG AGGTGCGGGC CAGAGGAGAC AGAGCCTAAC GCAGAGGAAG GGACCTGGCA	4200
GTCGGGAGCT GACTCCAGCC GGCGAAACCT ACAGCCCTCG CTCGAGAGAG CAGCGAGCTG	4260
GGCAGGAGCC TGGGACAGCA AAGCGCAGAG CAATCAGCAG AGCCGGCCCG GAGCTCCGTG	4320
CAACCGGCAA CTCGGATCCA CGAAGCCCAC CGAGCTCCCG CCGCCGGTCT GGGCAGC	4377
ATG AGA CGC GCG GCG CTC TGG CTC TGG CTC TGC GCG CTG GCG CTG CGC Met Arg Arg Ala Ala Leu Trp Leu Trp Leu Cys Ala Leu Ala Leu Arg 1 5 10 15	4425
CTG CAG CCT GCC CTC CCG GTGAGTGTGG CCCGGGGCAG GGCTGGGAGG Leu Gln Pro Ala Leu Pro 20	4473
CGGCGGGAAG CCGGGACTCG CCACTCGCCG ATGCCATGCA GGCGGCAGCA CGTGGAGGGG	4533
GAGGGGAGCG GGGACTTCTT CCCGCGCTGC CTGGCGGATC CTGGGATGGT GAGCCCTTTA	4593
ATGAGGACTC CTGTCCCAAT TCCTCTACGG TCCGTGGATG CCAGGAGGCT ATCCCAGCTC	4653
GTGGTCCGGG CGTCCTGCAG AGTGGAACCT CCATTGGTTC CCCGCTCCCA ATTAAGTAAA	4713
ACGACTCCAC AGGGGTCTGA GTCGCCGGCC TTAGGCGCTC CGCCGGCCTT AGGCGCCGCT	4773
TGGAGTTGCT CTCTCCCGTT GCTGTCTTGC TGGCCATCTC AGCGGCCTGG CCTCCGCCAG	4833
TGTCCCGGAG GATGCAGTGG CCATGGCCAA ACGCCTTTTC CATAGACCCT AATTCAAACC	4893
AGACTGCAGG CTGCACCCCC AGCGCCGCGG AGTCCGGGCG CTCGGCCCTT TGCACCGGGG	.4953
CAAGTTTGGG CACAGCAGAG CCGGCGCGGG AACAGGGGGA AGCTGACGTT CGGGGTGGCG	5013
GGAGGGACGG GATTAAGGCT GTTTGTGGGA CACAAGAGGG TGGCTCAGGG ACTTCGGTTT	5073
TTCTCTGGCT GCCCCAGGTG AGCCGGCCG AGCTGGCAGC GGGAGGTTCC GGGAAGTTGG	5133
CTTCAGAACG CTGAAGACCC TAAGAACCCA ACTTTGGGGT CGCTGAAGTT GTGCTGCCCC	5193
CGGAGGGCCT CCTCCGCATG GCCCGCGCGG GGGACCCTCC CCGCGAGTGG ACCCCGGTAC	5253
GGCTCTTCCC CTCCCCCGAC TCGGCTTTGT GCTGAAGCCG CGCGTAGGGA AGGCGGGTCC	5313
CTTGGCCCGC CCAGTAGGGC CGCGGGGAAA GAGGGACGAA CGTGGAGCTG GCGACTGGTG	5373
GGGGAAGCTT CTGGGTAGGA TGCAGCCATC CACCTTTGGT GGGGTCGGTC TCTCTAATCA	5433
GCGGCTTGGC GACAAAGAGC TTGGTCGAGG GTACCCCAGA AAGTGCTCTC CCGCCCCAAG	5493
CCGCCGTCGC TAGCCCGCCT TCCCAACGGG CGCTTTGTTC TCGGCCCCTG TAACCCTTCC	5553
CTGGGAACCG CCCCGCAGCG CTGGTCCTTG ACGTGGGCCG GGTCCTGGGT CGCCGCCAGT	5613
GTCAGCGCTG CCCTCCGGTG TCCACGCCCC TAGCCCCCGC ACCCGCTGTG AAGTCCCGGG	5673

1	TGTCCTTTCC	ACTGGCGCTT	TGCCCAACCC	CTGGAAGGCA	GAGGCGAGGT	GCGGAGCCTC	5733
	AGGCTTTATC	CTCCCGGAAG	TGGCAGTCTC	CCACCGCCAC	ATCTGGTCTG	CTTAACTTCG	5793
	ATAGTCCTGG	CAAAGGCAGA	CACGTGCACA	GGGAAGGAGA	GTTGAGCGCT	GGTAGATACC	5853
	AAGGTCGTGT	ACAAATAAAG	TGGCACACGA	CACGTCCCCA	GTCACTGTTA	ATGCATTGCC	5913
	TTCGCTCCTT	CCCAGGTGGC	TGGTGCTCTC	CATCACTCTG	GAGCCCAAGA	GAGGGCCTCC	5973
	ATAATTGTAT	TGCCCATGAG	TTGGGGTTGT	GTGGGGGCGC	CAAATCAGGG	TTCTCTGGGA	6033
	GGGCTATGAA	TTCCGAACTG	AGTCTCCTGT	GCACTCCTGG	CTTTAAGGTT	CAAGAAATTG	6093
	TTTGAGGGTT	GTGGTTTTTG	TGGGACTCAG	ATTATGCCTG	GAATCATAGT	TACCACTGTG	6153
	GAGAAGAAAG	TGGAGCTACT	TAGCATGCCT	CCCCGGCCCG	CCTGGCATTA	CCTCCGGCTC	6213
	TGTTCTCTAG	GCCCAACGTG	AGGCCTCACT	GGGGCAGTAC	AGATGCAGTA	CTGAATTTCT	6273
	TTCCAGCCAG	GATCTGGAGA	GGTGGTGTTC	TCTTCCCTGG	TGTCTTTAGA	GAGGCAGATA	6333
	TTCCTGTGAC	CTAAGCCCCT	CAAGCACCCA	TTAATAATGC	TGAGTAGACA	ACTAGAGGTG	6393
	GCGTTTTCCG	GAACTTCCTG	TGTGCTGGCC	TGGGAGGTTG	AACCCTCTAG	GAAACAGGTC	6453
	TAGGAAGTAG	AATTATCTCA	ATGGAAGGCT	TCCTGGAGGA	AGAAGATGAG	CTGAGCCCCC	6513
	AGGTCACTGT	CTGAGCTTTA	GGATCAGACT	CCCACTTGGA	GGCAAGAGTG	TTCGTTTTAC	6573
	TTTTTTTTT	TAAGTTTAGT	TTATTTTCTC	TCTAACAGAA	AACAAACAAA	CAAACAAAAA	6633
	AAAACCCCAC	ATTGTTTAAA	AGTGGGTGCA	TÄAGAGTGAG	GACATATTCA	GAGCTTCCCC	6693
	TTTTCCTGAA	AAATGAAGGC	AGCTGGGATT	TACTTAAAAT	GAGAGCACAT	ATCACAATTG	6753
	CCAGAGAGCT	GGTCCCTTTC	TCAGGGCTCC	CTAAGCTCCT	GTGGGAAGCA	GGTCAGACAG	6813
	CCCTGGGGAC	CAGAGAGATA	GGGAGTGCTT	TTGGGTGCCT	GCCTTTGAAT	GGGGAAGGGG	6873
	GGGGGAGCTG	CTGGGATCAG	AGGCTGCTAG	CAACTACTCC	CCAGAGACTG	AAGCAGGTTT	6933
	GTCCCTCAGT	GTCCTGTGGT	CTTCTGTTTC	TCCTATATAG	AATAGGAGAA	ATGGTTATTT	6993
	GCTCTGGAAT	AGTGACTTGC	TATTTGTTCC	CTTTCTTTCC	TCTCCCTTAC	TGTAATCATT	7053
	TGGACTAGTA	GAGACACTTT	CCCCAGGTCT	GGCAGAATGG	GAGGGAGTGG	GGGAGGCCTG	7113
	TGCTTGCATG	ATGTCACTGC	TGGCTTCAGC	TCTCCAGGGA	GGGTGGAGTT	GGTTGTAACC	7173
	TACCTGTGGC	TCTTGATGGG	CCACAATAAA	ACCTCATTAA	CACACATTGG	TAGGGAGAAG	7233
	GGACTGGAAA	GAATGATGGG	AAAGATTGAT	GTTTTTCCTT	TTTTTTTTT	TTTTTTTTG	7293
	GCAGTACTTT	CTAGATCTCC	CCTCCCCCTT	GCTGCAGCAA	AATTTTGGAT	TCCTGAAGTC	7353
	CTTTGAGAAT	GTATAATGGT	AGCCAGACTT	TTTTTTTTC	AGTCAGCTCA	AAATTGCCTC	7413

7473	GTTTTGTTTT	TTGTTGTTTT	GTTGTTGTTG	GTTTTTTGTT	ATCCTTGGTT	CTTATAAAGT
7533	GACCAGGCTG	TCAATATGTA	GCCCTGGAAC	AGTCCTGGCT	TTCTCTGTAT	AAGACAAGGT
7593	AAAGGTGTAG	TGCTGGGCCT	TAACTTTCAG	CACCTACTTC	CAAAGAAATC	GCCTCAAACT
7653	TCTGAAACCC	TGAGCAGGAT	AGTCTTACTT	TTTACAAAGC	GTGCTCAACT	GCCACCAAAA
7713	CCTCATGATG	GTATTTAGTC	ACTGCTAGGT	TCAACAATAC	TCTGTTATCT	TTATTTCCTT
7773	GCTCTGAAGA	GTTTTTGGTG	CAGTCTCCTG	CCAGGTCAAG	TCAAGTGGCG	CTGGGCCTCC
7833	TCTCAGTCTT	CTCTTTTCCT	TTCGGAGCTT	GCAGTTTGAA	CCAGTGACTG	AGACTGTGTC
7893	ATTCTAGTTT	TCTCTGTTTA	GCCTGGAGCT	GTGTGCCCAA	AGTGACACTG	TGGCAGGCAG
7953	AACACGAAGG	AATTCTTATA	GGTTGGTTAT	AAACAAATCA	TCAGACTGAA	ATTTTCTTTA
8013	ATCTCTGTTG	TCCCTCGGGA	GGGTCTGATG	CCGGCTGTGT	GCGTACGTCT	TCTCACCTTT
8073	AAGCGTGCTG	GAATGGACAG	GGGCAAGGTA	CGTGTAGAAA	AGTGTGTGTG	AGGCTGCTGC
8133	GCCTAGAGAA	CCGGTGAAGA	AAGCACTGGC	CTAAATGATG	TGTCCTGTTC	CCCACCCCAC
8193	GTTCCAGCTT	AGGAGCTTGA	GGAAGCACAC	CACAATGCCA	GGGAGATGCA	CTCCCTCGGT
8253	ACAAACAAGG	CCCTGGACTA	GCTCCAGCTG	GACTTTATCA	TCTCTTTGGT	GGCAGTGTCT
8313	CTGATCCTCA	CŢGTTTGAGA	GTCCTTGGTT	ATAATCGAAG	CTCAGTATTG	CTAGCTCACT
8373	CTGAAATTAC	TTTCAAAGAG	CTGTCTCAAC	AGCAATTCTC	TTGAACTCTT	CTCGGTAGCC
8433	TGAACGACTC	TCCTTGACTG	TTGTTCCTAA	GACTGAAACC	CACCATATGC	AGACTCGAGC
8493	CTACATAATC	AGTTCGCGTC	TGTATGTTTT	ATTTCTTTAG	TTCTTTCTCC	TTGGGTTTGG
8553	GAGCCTCACA	GGGTCCAGCA	AGACAGCATT	AACAGGTTAG	ACTTAGAAAC	TATTGCCCAT
8613	TTCCAACTCC	AGTGACTCAC	TGTCAGCTCA	TGATTTACCG	GTCCTGCCAC	CTGAAGCTCA
8673	ACATTCTGTC	CTTTCTGCCC	CATACCTGCT	TAGACATCAC	ATCTGTAGAG	TCTGCTCCCC
8733	GTGCTGGGGA	TGTAAGTAAA	AAAAGTGCTT	ACGATGGTGC	TCATTTCATA	ATTAACATGT.
8793	TGTGTGTGG	GAGTGCCTGT	ACTTTTTATT	GTTAGGGTTA	GTCGATAATG	AATGTTAGCT
8853	CTTAGCTTTT	TTCTTTCCTA	GTTTTCTTAC	AGGCTTGGTA	GTTTTTTAG	GTTGGGTGGG
8913	CACTGAGGCA	TGAGTGTGTG	CCTGATTGTT	TGTATCATTG	CTTTATGGTA	CTTCCTAAGC
8973	TGTATGGTGT	TGCTCACATA	CGTGCTCTCG	TATGCTTGTG	TGTTTGAGAG	CGCCTGTGCA
9033	GAGCCCTGCC	TGAATCCTGT	TGGGGÇTGGC	GCCGGCACAC	TAGAGTGCAG	GAATACACTG
9093	GGAGTGACTG	CATTTTGTGT	TGCTTGTGAG	TGGACACTCC	AGATCTTCCT	TGGAGTTTGC
9153	GGAAACACCC	CTGAGTATTG	TGGGTAAACC	ATTGTGCCTT	TGTAGCCTAC	TTTAGCTGGC

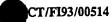


maccomcmcC	CHCHCHCHCC	СССАСССТТС	CTTGGGTACA	GCTAAGAACT	CTTCATAGAA	9213
						9273
•			GAGTGCTAAG			
CTGCTTGCCC	TCTCATGCAG	TTTATCTTGA	GCTTGGCGAA	CACACTTACA	GATTTAGTAG	9333
AGCTTTTGTC	AGCCCTGGGA	GGTGGGTTTC	GTGGCCACAA	GTGGGTAGCT	TGGAATCCAA	9393
GACTCCTGGC	TTCTAGGTTG	CATTCTCCTG	TGGTTCTTTC	CAAGGGAATG	CTAGGGGAAC	9453
ATTTTGGACA	TTAGATTATT	TCTAGTCCCA	AAGCACACAG	AACATACTGT	TTCCTAATTG	9513
CCTTTTTTT	GTTTTCCTCT	CAATCTGGTT	TTGAAGTGTT	GGGTTTGAAA	ATTGCCCCCT	9573
GAGAGCCTGC	CCTAGTGTGT	GCAGAGGGAA	GATAGTGGAA	CAGGAAGTCT	GTAGAAAGTA	9633
TCTTCCTTTC	CAGGACCTTG	TGCCCCGGAG	CAGAGTCAGC	ATGGTGTCAT	ATCGCTTTTG	9693
GCTATTCCAG	AAGAGATGAG	GTTTTAGGTG	AGAATGAACC	TTTTAGAACC	TTCTAGAACC	9753
TTCTGTTGAG	TATGACAGGA	ATGCCCTGAA	TAGGGTCCGA	AGTGCATGGC	CACTTGTTTG	9813
TCTTTTCCAT	AAGCAAGCAG	CTTCAGGTAC	AGACAATAAG	ACTAGGTTCT	TGGAGTGAGA	9873
CCCTGCACTT	GGTGCCATTT	CAGCTCCAGA	TGGACACTGG	AGGTCCCTAC	ACAGCAGGCT	9933
CTGGGATGGC	TGGCTTTGCT	ATGTACTGTT	GCCTGCTCTA	CAAGAGCTTC	CCAGGTTACT	9993
AGCCTTTGTC	GACGCTGGGC	TCGCTGGCCA	GGCTTGGGCA	TTGGAGAAGG	GACAACTTGC	10053
CACCTGGCAT	AGGCTGTGTG	TTTGGAGAGT	CAGGAGGTCT	GGTGAAGCCC	GCAAGTGGAG	10113
GCAAGTTTAG	TGGGACTTGA	GGAGAGCTCA	GTAGGAAATC	TCTGGGCTAG	TGACAGGCAG	10173
GTGTGGTGGT	GGTGGCGAGG	TGGCGGGTCT	AGATCTCCTT	TTAGAGATTT	GCCTAGGGAT	10233
CGTCCCTGCT	GACTCTGGAA	ÇTCAGAGGCC	TCCAGAGGTG	TCTCCTCTGG	GAGCCTCTCA	10293
AGGGTCTCCC	ATCTCCTACT	GTTTATGGCT	TTGTGGGCTA	CCTAATTACA	TAGAGAAGAT	10353
ATGTTCCTCT	GCCTCCAGCC	CTGGAAAGTT	CTGCCCAGTG	ACTCACCTGA	GCCTGCAGCC	10413
ATGTGTGTAC	ACAGGCGCTC	TCAGGGGCTT	CTGTCCTGCT	GGCTTCAGCC	TTTCTAGCCC	10473
CTGGTGTTCT	CGGCAGTGGT	AGCATCTGGG	AAACCGGGTC	ACCTCTTATT	TGCAGCTCCC	10533
TCCCTTTCTT	GGTGTCTTCC	CCCTTTTTAA	CTACTGGTCT	GATGGCCTTA	GACTCATGCT	10593
GAAATTCTCC	TTTCTTTTGT	CCTAGCCTTG	TCTCTGACTT	CTTGTGATCC	TCTGGGCCTG	10653
TGAAATCCGC	TCAGGGGCCT	CCATTTCTAA	CAGTCACACA	CTGGTGGAGA	GACCGAGTCC	10713
TĢGGATGGTG	AAGCTAACCC	TGCTGGGCTT	CTCAAGCTTC	ATTTGGTTTC	TCTTTATTCC	10773
TTCTGGAGGT	ACTGCCTGCC	CCAGGGGAGT	CTCAGACTAG	ACCACTCTGG	AGTTGGAGGT	10833
GGGGCAGGTT	TTCAGATCAG	TGCCCTTGGC	ATTCGTTGTG	GGAATGGGGT	GGATGGGGCC	10893

•		•			•	
TCTGGGCAAG	GTCAGGCTGG	GGGTGGAGGC	CAGGTGATGT	TCTCCGCACC	CACACCCAGG	10953
CAGCCTGGCA	CCCTCCCCAA	GGTCCGCTCA	TCAGCAGGAA	TGAAAGCAGT	GCCGGGCAGG	11013
TTGGGGCAGT	GGGCAGGTGG	GCGTGTTTAT	CGCTGTGCTC	ATCAGCTGAG	TCACGATGCC	11073
AGGCCCCACA	AGTCCTCCCT	GGAGGCTCAC	CCCACCCACC	TTGACCCACC	AGCACCCACT	11133
AGCAGGAGGT	AGGGCAGGGC	AGTGAGACAA	GACCAGCCTG	GGGGTCTGAG	AGGCAAAGGG	11193
GAGTTGTTCA	TGACCTGGCT	GTGCATGGGG	ACTTGTGGGT	GTCTCAGATA	TCTCTGCTGT	11253
CCAGGAGGAA	GCTGTCTTAA	GTGCCAACCT	GCCTAGAGCC	CCTGCTGGGT	GCAGGAAATG	11313
CACAAGGGAG	AGTGCCCATC	CATGGAATAG	GCCCATGGAG	CTAGACCAGT	GACAGTGACA	11373
GTGAAGTCAG	CCCCCACCTG	TGTCTTCCGA	GCCAGCTGGA	GGGTTTTTAT	CTCAGATTCT	11433
GCGAAACCAT	AGAATCTAGT	CAGGAGCCTA	GACTGCAAAG	CAGGCTTCGT	TGATGCTTTA	11493
ACTTGCAGGC	TTCCTGGGTA	TGAGGGATAC	TTAGAAAGGT	CCCGCAGGTA	GGGAGGGCAT	11553
CAGGAAGTAG	AAGAGGGCCA	GGCACTTCTA	TCTCCTGCAT	TGCCCCCTTC	TCCCATCTCC	11613
AAGGATGGTA	AAAAGAACCC	TTCCAGTACA	CTGACAGAGA	GGAAAACCCT	TCATCTCACC	11673
CCATTTGGAT	CTGTCGTATC	AGCATGTGCT	GGCCCTGCTT	CCATACCAGA	GGTGGCTAGA	11733
GATGTTCCCT	GGGAATTCAC	TGGTTGGGGA	CTTGAGTGTA	TCAGAGGGGC	ACAAAGTAAC	11793
ATTAACTCTG	GTATCCTCTG	CAGCAAATCG	GAGATCCCCT	CTCCTAGGCG	AGTTCTCAGT	11853
GGATATGGAG	GTCAGGTTTG	GGCTTGTAGG	GCCCCAGCAA	GAGTCGTTGA	TGTCACTCCA	11913
GCTTCTCCCG	AGGAAGATGA	GGGTGCTGTG	TTGGGATCAC	ATCTCTCCCT	GAATGGCATG	11973
TTGGGGAGGG	ATGGAGCCCT	TGCTTCTGAC	CCCTAAGCTT	GGTCTTTAGG	TGGCCACAGT	12033
CTCTGGGTTC	TGTCCTACCT	CCCTGCCCTT	GTGTGCTTCA	AAGGCATGCT	AAAGGGACTC	12093
TCGGCCATTC	CGAATGGCAC	AGTGTTCCTT	CTGTTCTCCC	ACCCCCAGAA	GGAGGCAGGC	12153
CTGGATTGTA	GATTCCTAGA	AGTAAGTGGC	CCTGAGCATG	CTGTTGATGA	ACCTGGAACC	12213
AGGCAGGCTG	GGCATCCTAG	GACCTGTCTT	TCCATAGAAG	TCTGAATCAG	TCTACCTTTG	12273
GGACTGAGTA	AGGGGCTCCT	CACATATCAG	CTGGCTAGTC	CATCTTGGCT	GATCTAAACC	12333
ACATTAGGCT	GAAGAGAAGC	ATGGTGTACA	GTCTGGTCCA	CCCGAACCAC	ATACTGGCTT	12393
TATCAGTTCT	CGTATAATTT	TGCAGGTAAC	TTTTTAGCTC	TAAGCCTGTC	TCCTCATCTG	12453
TGAAATCGGG	TCCCTCATAT	CCTGCCTAGA	AGGGCTTTTG	AAAAGATTAA	TGAAGTAGTA	12513
TGCCGAGTGG	TTGGGGTTCT	CTCCTTGACT	GGAGCAAGTC	TCTAGGAGTA	CTAAGGATAG	12573
CCTGCTGTGT	GCAGCACCCC	CAGGGACTGT	GCCTGAGTAG	GAGGGTACAG	AGTCTTCATG	12633

TGAATGGCCC	TTCTGGTCTT	GCCCGAAGT	TAGTGTTGAT	GTCATAGAGT	CTACAAACAT	12693
GCCTTTTGTC	CTTCCTCAGA	AGTCCAAGCC	TTTCCTGGCA	GACCAGACAT	TCATCTCCAC	12753
TGAGCCTCTA	TGTGAGACTG	GCTCCTGGCC	TGAGCTGTGT	GGGCTGAGCT	GGCGAATGGG	12813
AAAACTAGAC	ACCTGGGCAC	CTGGGTGGGG	GCTCGGGACA	GCAGTGTTTC	AĢTTGTAGGC	12873
ACTGTGCCCC	TGCCTGGAGC	TTCTGACTGA	AGGTTACCCT	GAGAGGAAGC	AGGTTCCCTA	12933
TAGACACTAA	CATAGCTGGG	TCAGAGTGCA	AGGTGGGTGT	GCCCTGCCC	TGACCCATTC	12993
AGTGCAAAGG	CTGCTCTTCT	GGGAGTGAGA	GCTCTGACAG	GACTGTGATG	GCCGAGGGGT	13053
CTCAGAGCAA	ACCTGCCTGG	CCTCTCCCCA	CTCTGATGGA	TATGTGCTCT	TAAACAAGTG	13113
ACTGTCCACT	TTGCCTCAAT	TTCAACATCT	GTAAGATAGA	TAGGGCGTTA	TGGTCTGAAA	13173
ATGGTTTTAA	AGATTAGTTA	GCTAATACAG	GGAAAGTGCT	CTGACAGGTA	CCTGGCACCT	13233
TACTCAACAA	GTGGCTGGAG	TGCCTGATTT	CCTAAGGTCT	CGACCTGTCC	CTATGCTTCA	13293
AGTGCCCCTA	CAGCCTTGGT	CAGGCCCTTA	GGTTCTCCCA	CCCACCGCTG	GCCCCAGGAC	13353
CTAGACTGCT	GGACCCTGAC	CCCATTTTTC	CTTTAAGCCA	CCTCTGCGTC	AACTCTAAAA	13413
GGCGGTGGAG	TTGTTTATCT	AGGCTGTGAG	GTGTCAGAGA	AAGGACCTGG	GCCGCTTTGT	13473
TCCTGTGTGG	GCTGGGGCCA	CTCCAGGAAC	TGAGAAACCC	ACCCACCTTT	TCAAAAACAG	13533
CCTCTTCTCA	GAGTCTGGCA	CCTCAGCTAG	CCACCATGCT	GTGGGACCAC	TCCCAGCATG	13593
CTCTGCCTTT	GGTTTGTTTC	CCAGGGGCCT	CAGTGCCTTT	TAAAGATGCA	CAGGCATCTT	13653
TAGTTCAAGG	GGAAAGAGGA	AATGAAGTGT	ATTTGCTGGT	GGTGGTATTC	CTGTCACTTG	13713
CATTCTCACA	GAGGCTAAAG	AAATTTGCTC	TTTGTATCTT	CTAGTCTCTT	CTTTATGATC	13773
TTTTCCCATC	TGTTGTATCC	CAACTGCAGG	GCCCCAGTTC	TAGAATTAGC	CCCTCCCCA	13833
TAGGAAGCCG	ACTTATGCTA	TAATGTGAAT	GACAAGTATC	CTTTAGCCCT	TCCCACAGGC	13893
ATTTTAATTT	TCAAAAGGGC	ATTGCACAAC	CGCAGAGACA	CTAAGAAGAG	AGGTTTGGTG	13953
ATCAGAGTTA	CAGCCCCAGC	CTCCCAGCTG	GTGGCCCGGC	TGGTGCAGGT	GTGTCGAAAG	14013
CAGTAGTTTC	TGCTTCAGTG	AAACTTGAGG	ATCCTTTATT	TAGCCAGTTC	AGGGGCGGAA	14073
TGGCCATGCG	AAGTCTATGT	GTCACAGGTG	TCAGGCCCCC	ATATCCTGCT	GAGTCTAGAA	14133
TCAGCTACGT	AGCAGTTTTG	GGGGTATTGC	CAGACTGGGA	GTTTACATCC	CAGAAGCGAG	14193
AATGGTGGGG	TTCCTATACT	GCTCCAGACA	GGATCTTTCC	CCCAAGTTTG	TCAGCCACCT	14253
CTCTTCAAGT	CCCTTGGCTC	TGACCAGCAA	GACGTATCCA	AAAGAAACTG	AGGAGGCCCT	14313
TCACTTCTTT	TTAGGATAGT	GTGGGGCCAG	CATGGTGGGG	GTTGGGAATG	GCTTTCTGTC	14373

TCTTCCATCA	TCACAGGCTA	CTTCCCAGAG	ACACTTTGAT	TCTGGGCATC	TCCAGCAGTC	14433
ACCTGGCCCA	CAATGCTTTG	CTGCCCTTTG	CTTCAGCCAC	TGTATCTGGT	TGTCCCTTGA	14493
AGGTGAGCCA	GAGCTCCTAG	GCAGAGAGCA	TGTGCTATAC	AAAGCCGTAG	GCTGGGCCCT	14553
GGGAACCTTC	TTGCTGTCAT	CCTCCTGTCA	AACCCCTATG	GTATGGTAGC	CCACATAAGG	14613
CTTGTGCAAA	AAACAGGCCA	AAACATAAGT	TATCTTTTCA	CTCTATCGGG	TCTTCTCATT	14673
TTCCCATGGT	ACGTTCGGCT	GGCCAGGCCC	AAAAGATTTG	AAGAGAGGTG	GCTGGCAAGT	14733
CTAGGGGAAT	AGGTCTATCT	GGTTCCCTCC	AGGAGCAGTG	CCTAGTGAGA	GGCTGGGCTG	14793
GGCAGGGCAG	GGCCCCTTGC	TCCACATTGC	CTGAAGTCCC	GCCCTGCCCG	TCCTGGCTGG	14853
GATCTGGCAG	GTCTTCCAGC	TCCACACCCG	GCTCTCAGCT	GAGCCTGCTC	AGAGACTAGT	14913
CCTGGCATGT	GGGTTGCAGG	GCTGGTTCCA	GCTCCACCAG	GAGGTATGGG	CGTCTGGGTA	14973
CTCATGGGAC	ATTGACCTGT	AGTGGGTATG	GAGAGTGGAG	GAATGGTACA	GGCAGGTGTG	15033
CTGGTGCTGA	CGGACTTGAC	TCCGGCATTG	ACCTTGGCTT	GCAGTCTGGT	GTTAAACTAA	15093
CAGGGAATGC	TGACAAAAA	GACAGTTATT	AAAACCAAGA	CAGGATACTG	CTTTCCCACT	15153
CAGCCCATTC	CCAAGAATCC	CCAAGACGTA	CAGGAAATGT	GCAACAGCAG	TGGGAATTGC	15213
TGAGTTGGGG	GATGTGGGTG	AGCTGTGTGC	TCCCAGGGAA	TTTTGGGAAA	TTCCCCTCCG	15273
TTGAAATGCT	GTCAGGGTCT	GAGCCTTGGA	GGTGTTTTTG	GGGTGCTGTG	CTCCCCAGCT	15333
AAGCAGCTAA	CAGTCCTCTT	TACCTGCCTT	GTCCTCACCT	TGCCCCACCC	TGGGTTGGGC	15393
CTCTCGTTCA	CTCCCTGCTG	GGTCACCAGT	ACTTCAGTGC	AGGTCTCAGC	TTGATTCTTG	15453
GTGGAGAGAG	AGAAAGTTGA	TAAATCAGGG	TGCCTGTCAG	CCGGAAATTT	GGGTGTGTCC	15513
TGAAGGCACC	AATGGGGGCC	CTCCCTTCTG	GAGGTGGCTT	TAGGAAGGGG	TTTCTGGGTC	15573
TTGAGGCCTC	CTTACAGTTT	CTTAGCTCCA	TGGGAGAGAA	GTGAGGAGTT	GGGTATCGTC	15633
ACCCCAGCAT	GAATCTCTGG	TCACCTCTCA	GCATGCACTG	TCCAGCCTGA	TCTTTGAGTG	15693
CCAȚAAAAGA	ACAGAATTAT	CCTCTCAGAG	CACTTCATTT	CCCGCCAGCA	CAGTGGGTAC	15753
AGAGACAAGC	TGCCCAGACT	CCCAGCGAGG	GACTAGTTGA	GCCCCAGCAT	GGGACTAGTT	15813
GAGCTAGACC	TGATACAGTC	CCAGAGAGCC	TCGTTGAGGA	AGCTTTGGGA	AAATTCACCC	15873
AGCATTTCAG	CCAGGACTGG	AGGAAAAGGT	GATTATGGGA	AAGAGAGCAG	TCAAGACCCC	15933
AGGCTGTAGG	ACACAGGATA	CAAACTGAGA	GCTACCGGAT	AGGAGTAGGT	TTTAGTCACA	15993
ATCTCTCCTG	TCCGCCCTAC	CCTCCAGGAG	ACATTGCACC	TTGTAGAACA	GCTGCCCCGG	16053
AGTCCACCTT	TGGGCCCCCC	TGGGTAGCTC	AGTAGTGTCA	GCATCCTCTC	ATTGACATCA	16113



	GTCAGGTTAC	ACAGTGGGGC	AGCTAATGTG	AAGGCGCTAG	GCTGGGAAGC	CAGCTACTTG	16173
	GGAAAACTAG	GTTGTTCCTG	GTAGGCCCTA	GCAGGAAGGC	AGTTCCTCCT	TTTCTTGGTG	16233
	GCTTTAGGGG	TCTTTGGAAG	CTTTGAATGT	TCCCTCAGCT	CGTTGGTGAA	GCAGGCCCTC	16293
	CTGGTACTGT	GGTGTTTGTC	TTCGAAGAGT	GAAGGCATTG	GAAGTAAAGA	CTGATGGGGC	16353
•	GCCTTCCCAG	GATGCTTTGC	TTCTTGCGCT	GGCTTACAGA	GCTCTCTTGC	TACCTAGTGC	16413
	CTTGACTTTG	AACACCAGAT	TCAGTCAGGG	AACAGGAGTA	GAGGTCTTGC	CTTGCTGAGC	16473
	CCCTGCGCAC	TGCAGGAAAA	GACTCCTCTG	AGTGGAGCCT	TTCCTCCTCA	GGTGACTGCT	16533
	TTCAAAGTAC	AGCAGCCTCT	GAGGGGGAAG	TGTCATTTGA	CATTGTGGTA	GTTCTTGGGG	16593
	TCCCTGGATA	CAGATGTCAT	GCCCAGATCA	TAGGTCTGTT	TGTACAGAGG	GAGGCGAGTT	16653
	CTGTAGCTCA	GAGTCCTCAG	TACCCCAGAG	TTGTGGCTCT	AGGGGTGAGA	GGAGAAGACT	16713
	ACAGCCCTTC	AATCACAGGT	CTGACCTGTG	GGTAGGGGTA	GATCTCTTGC	ATACTATGAA	16773
	CCTGTTTGAA	ACCCCTGGGT	ATTTGCTGTG	GAATAGAGTC	TTGGTTGGGT	AAGAATGGTG	16833
	GATGTTTATC	TTGGTGTGAC	TCTCGGGTGG	GGGTGGGGGA	TATGTCCCTG	TCTTTCCCAA	16893
	TGTAGTATGC	TGAGTGGACA	GAGACCGTGT	GACTGAAGCC	TGGGCTCCTG	GAACAGGTGT	16953
	GTGTTGGTGG	GGGGTGGGGC	GCAACTATCT	GGGATCCAGA	CTGCTTGGGA	ATGGCTGTGA	17013
	CCCAGCTCCT	TTGATAACAG	CAGCTCTTTG	TCACTGGATG	TTGTGACTAA	TGGGACTTGT	17073
	TGATTCAGTT	ACTCGGCTCC	CACCCACAGA	CGCCGGGGCT	TCTGTTGTGG	CACCAGGCAG	17133
	CTGCAGACGG	CCCACAAGTT	TGCCTCGCTT	TCCCACTCCA	CGAAGGTAAG	TTCCCAGCAC	17193
	TGCCCAAATT	AGAGACTTGT	GAGTGGTCCC	CTCATACCCC	ACTCCCTGAG	GCTTCTCCTG	17253
	GAAGGCCTGG	AATGGGGCAC	TGGGTGTGTA	CGTGCTGTGG	TTTCTGTTAG	GGTCAAGACC	17313
	AGGCTGTTTC	TTACCTGGCT	CGTACCTCCA	AGTTTCCAGG	TGATGAGTCC	TGATTTTTGA	17373
	AGTGAAGGAA	TCCATTTAAT	ATCAAAATTC	TGTGACCTTA	AATTTTTTTC	TTTTATTATG	17433
	TGTCATTTCA	TATGTACGCA	TATTTTTTG	TCTGTGTGTG	GACATGCTTG	TGGCGATCAG	17493
	AGGACACTTC	AGAAAGTCAG	TTCTCTCCTG	CCGTGTGGGT	CCTGGGGAAT	CAAATCCAAG	17553
	TTGTCAGGCT	TTATCCTGAA	AATAAAAAGT	AGACAGCCCT	TGGGATCCAA	AGCTTCTTAG	17613
	GGCTGTGTGT	CTTAGACACC	ACCAGTGTTG	CACAGCTGGT	AACATGACAG	TGTCCTGGAG	17673
	TGCTGATTGG	AAGCCACAGG	CCTCTGTGCA	GGGCGGTAGA	CTTCCAGGGT	ACGGGGCAGG	17733
	TGGGCGTTCT	СТАСАААААС	CTTGTAATCG	CGGACGTCTT	GGAGATGCCC	CCTAGGTATC	17793
	ATGATTTTGG	TGTGTGACAC	AGCTGAACTG	TCTTCATACT	CAGGATATCA	TGAAGTGCTG	17853

					•	
GGGTGCAGAC	CACTCTCAGC	CTCAGGCAGC	CAGGACCCGG	GGCTCCATCA	GATTGCGGTG	17913
ACTACCACAG	AGGGTGGCCT	TCCTTCCGGT	CAGTGTGGGT	GTGGGAGCTG	GCAGGAAGTG	17973
GCTCCAGGCT	TCCTTTAAGC	ATCCTCTGCC	CACAGCCCCA	AACATGTTCT	TTGGCAATGG	18033
CTTGCAACTA	GAGGTGAACT	CTCTCCTGTA	CTATGTCCTG	ACCCACGCTG	CTGCATCTAT	18093
TATACCTTTC	ACACGCGTGA	TGGGTACCCA	GCGGGGCTGC	TAGGCAGGGT	TAAGCACTCA	18153
TCTTGTTTCC	TGGTGCTGAA	GCTGTGGTAA	AGAAACTGAG	GCCATTTTCC	CTTGAGAGAG	18213
ATGGTCTCAG	CCAGGTCTTT	CTCGGCCTGG	GGAGCCCGGA	AGAAAGGATG	TACTACAGTG	18273
AGTGGACACT	TGTTGGCTGA	TGGCCTTGGT	AGGTCCTTCA	CCCTGGGAAG	TGCTGTTTCT	18333
TATCTGTTAG	AGATGCTGAC	CTCAGCAGGA	CTGGAGGAAC	TGCATGGGAG	GTGTAGGAAT	18393
GAAAGTGAGT	GGGGAAAATT	ATCTCCAGCC	CTAGGGAAGT	CTGAGGCCTG	TGTCCCCTTT	18453
GTCCTGGACT	GGGCCCCTGC	CTTGGGTGTC	TGTCCAGGGT	CTTTGCTCTA	CAGCCCCAGC	18513
GGATGCCCAA	AGTAGACGAG	TCAACTGGTC	CTTTCTTTCA	CCCTGTGTCC	ACTTCTCATG	18573
TATCTACCTT	CATAATCCTT	CTAGGTAAAA	CAAGCCTCTA	ACTTTGGGTT	TTCAAATCAG	18633
CCAGCTTCCA	GGCTCGATAG	TACGAACCAT	GAAAATCTTT	CTTACCATGA	GGTTGTTTTC	18693
TAGTGTGTGT	GTGTGTGTGT	GTGTGTGTGT	GTGTGTGTGT	GTGTGTGTAC	GTACACATAT	18753
GTACCTCTAT	CAGTGTGCTG	TGCGTGTACC	ACAGCAGACT	CGTGAGGAGG	TCAGGCAAAC	18813
TTTATAAAAA	TCTTTTTTT	TTGCTTCACT	TGAGTCCCAG	GGTCACACAG	TGGCAAGTGC	18873
TGAGCTCTGT	TCTCTGTTCT	TGATTTGTTT	TGTGAGCAGC	TGATGTTCTT	AAGGCTTGCG	18933
GAGGGGAAAG	GTAGGGCTGG	CTTGCTTCTT	CCCCGAGTGG	CGGTCAATCC	CTAGACATCT	18993
CTAAGCCGTG	GCCACACGTC	CTGGAAGGAC	CCAGGTCAGA	AGTGATACTG	AGATGGCCCT	19053
GTGAGCCCTC	TCGAACACAC	AGGGTTGTAA	ATAGTACCTG	ATTGTTACAT	TGGAGACTCG	19113
TCAGCTGGGT	GGAGTCCTGG	TTCAGAGGGA	GTTATTCCTC	CCCCACATT	TCTTCTCTTC	19173
TGGGGCTGAA	GTCTCTTCCT	TCCTTACCTG	TGATGCTGTC	ATGATAGGTC	CCAGCTGAGA	19233
GTGGAGGCGG	GGCAGTCAGG	GAGCTGCTTC	TCTTTGCTTA	GCAGGGGTTG	GAGACTTGGG	19293
GTGTAGGGGT	TGGCTCCCCC	TTTCCCTGCC	CTGGACCTGG	TTTCTGGTTT	CAGCAGAGAT	19353
TCGTTCTAGA	AACTTGTTGC	GTAAACAAGA	TCACAAAGCG	ATAAGCTTGA	GCAAAACCCA	19413
GGGGAACAAA	TTGCTTCCCT	GTGAAGACCC	AATCTTAGCT	CTTAGAGAAG	CCCTCCCTTT	19473
TGGAAATTGC	TGACTTTCAG	GGCTTCTCTG	TGGAGGAAAG	AGGCTAGCCG	CCGTATGTTT	19533
GCCTGGATTC	CAATAAATCT	TTGCGGCCTT	GGCTACCCCT	TGTTGAACAA	GGTCTGCACT	19593

CCTAATGCGT	GCCTCAGGTG	GTCTGAGACC	TCTACCCCAT	CTCCAGCTTT	TCCTTCCTAT	19653
GGAGGGAGTC	AGTGGGTTAG	GAGAGAATGG	AGTTGAGTCC	TGGAATGAGG	AGGAAGCTAT	19713
GAACTCGGGG	CCTGTTCCTG	TCTGGTGGGT	GCTCTTCTCC	GCCGCTGAAG	GAGGCAGCCG	19773
CAGGGAAGAC	TACCACAGGA	ATCCGAGTAC	CACCTGGAGC	AGTGTATACA	GGATGTGGGC	19833
	TAAGGGCATG			•		19893
TGTGGATGGG	CCACAGGGAA	ATTTTTGAGT	GTCTACTGCA	GTAGTTCTCA	ACCTGTGGGT	19953
TGTGCGCCCC	TTGGTGGGAG	TTACATATTA	GATATTTACA	TTATGATTCA	TAACTGTAGC	20013
ААААТТАСАА	TTGTGAAAGA	ACCAAGAAAT	CACCGCAGCA	TGAGAACCTG	TATTAAAGGG	20073
TCACGGTGTT	AGGAGGGTTG	AGAGCCACTC	ATCCTCTGGG	TCTAGGCCAT	GGCGGGCTGT	20133
AACTGCTCTC	TGGAGTTAAG	CCACAGTGAA	CCAGCTGTCC	TTGCAGATGG	ACTTGTGGAG	20193
GCTCCAAACC	TTTGTCCCAG	GGGAGAAGAG	CTTGCTTTTG	CTTTGTACTT	TTAAAGGAAG	20253
TTCAGTGGTC	TTCGGGCCTT	GTGGCTGCTG	TGTGTGGAAG	TGCCCCTGTA	CAATAAGCTG	20313
TATAGATCGT	GTACAACTGC	AGTTTTCCTC	CGTGGGŤCCA	CCAACCACTC	CTGACTCCAC	20373
GGATGAGTGA	GGCCAGTAGG	GCTGTGTGTG	GGTCCCTAGG	CCAAGCATCC	TGGACCACGA	20433
TGAGCCTCAG	CTAGACCACT	CTGGATCTTT	AGCAGAGGCT	CCTAGAGAGC	TGGCTGGCTT	20493
CCTCCTGCCT	TCTTTTCTCT	TAAAACTTCG	TCTCAATCGG	AAGCTCCTCT	GTGCACGTGA	20553
CCTCCAGGCC	TGGGGGTCGC	CACAAATCCC	CTCATCACAA	GACGAGCAGC	TCGCATGAGG	20613
GACACGACAC	TTGTTACCTA	CCAGGCTGTG	GGGTTTTTGT	TGGTTGGTTG	TTTTGTTTTG	20673
TTTTGTTTT	TTACTTGTAC	: AGAAGTGTTG	TGACATCAGA	TGTCAGCTGT	TAGTGCTGGC	20733
ACCATTTTAC	AGGTAGGGAA	CTGAGGCTGT	AAGATGTGTA	GTGACATCGC	TAAGGCCACT	20793
CAGTTGGTG	GGCCTTACCA	AGGTCAGGTC	TTTGGAGCCT	TTTGCTGAAC	CATGTACTTC	20853
TATCTCTGT	TTGTTGAAAC	: AAAGTCTATA	TGGCTCTGGC	TAGCCTATAA	CCCCATATGT	20913
AGACGAGGC'	GACCTCGAAT	ACACTGCAGT	CTTTTATGTC	TGCCTTCTG	GTGGCAGGAT	20973
TGAAGGCAT	TGATTCCTCC	TAACTGTACA	CTTTAAAAAA	AAAATCATTO	TTTGTTCTGG	21033
TCTGTGCCA	GGCCTTGTA	A GATGTTCTGT	GCTGAGCTGG	GCTATTTGG	TTAGTCTCAT	21093
TGCTGAGCA	G GGCCCCTGT?	A TCTTCCTTCT	CTGTCACTTG	CTTACCTGG	TCTTCCTCCT	21153
GCACTAGCT	A TCCTAGAACO	AGTACTGAGA	GCAACTATGO	GCCCAACTC	GCCCCTTGCC	21213
CAGCCTGCT	r agctggggg	GGTGTTCCAC	TTCCCTGCC	AAGTCCTGT(G GGACTGTGTT	21273
TGTACTCCA	C CACCTTCAG	TCCTTGGAG	TGGAGCAGG	CAGGCGGCT	G CATTCCTGCA	21333

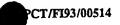
•	
GCTGCTGTTG CCAGGGAGAG CCCATCCCAT TCACTTCAGT CTCCTTAATG TAGAAGCCTT	21393
GTCGAATTAG CTTCCACTGT CCCCAACCCA AGAGTACCCT GTCCTTTCTT CACTAAGAAG	21453
GCCAGGATAC AGTCCTTCCT GTGGCTGATA AGACAGGCCT TGGGACAAGG CCTGGGACCA	21513
CACTGTGTGG GCAAAGCTGC TTCAGCACCG ATGGCTCCTC CATGCCAAGC TTGGCTCTGC	21573
TTCTCACAGT TGAGACTTCT GTGCGCACAC CCACTGTCTA GCTCAGCTGG ACACTGATTT	21633
TCTTTAAATG TATAGATTTT GGGGTGGGGT GTGCTGAAAG CTCCCACTGA TGCCCCAAGC	21693
CTGAGTCTCA GAGTATGATC AATTGATGGC TTTCATGGGT ATCACAGCTT CTGTTCCCAG	21753
GTCAGACTCC CTGACCAGTC AGAGCATCCT GGGGTTAGAC AATGTCCCCG TCACTTGTGC	21813
CTCCACCTGG CACCAGGCTA TGATGTTATG GCATTGAGGG TATGAGAAGG ACCAGGGGTT	21873
TCCCAGAGTT ACGCCCAGGC GCACAGGCAA TTGTTTCCTA CATGTGTGC TGGAATGGTT	21933
GGGTGAGCCT TTTCAGCTGC CTACAATAGG AACCCAGGGA TACTGGGCAT TGACCAAGGC	21993
ATATCTCATA CCCTTTTCTT ATCTTTCTGC AG CAA ATT GTG GCT GTA AAT GTT Gln Ile Val Ala Val Asn Val 25	22046
CCT CCT GAA GAT CAG GAT GGC TCT GGG GAT GAC TCT GAC AAC TTC TCT Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 35 40 45	22094
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser	22094 22146
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 35 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG	
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 35 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG Gly Ser Gly Thr	22146
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 35 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG Gly Ser Gly Thr GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC	22146
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 35 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG Gly Ser Gly Thr GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC	22146 22206 22266 22326
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 35 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG Gly Ser Gly Thr GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC ACCAGCCTAA GAGAAACTCT CAGAGAAATG AAGGGGTTCC ACCAGGCCAT GGGCAGCCAC	22146 22206 22266 22326 22386
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 35 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG Gly Ser Gly Thr GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC ACTCAGCTAA GAGAAACTCT CAGAGAAATG AAGGGGTTCC ACCAGGCCAT GGGCAGCCAC TAGAGCCAAC CCTTGGAGGA GTTTGACTCC ACTGAGCCTT GGTGTGGTGT	22146 22206 22266 22326 22386
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 35 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG Gly Ser Gly Thr GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC ACTCAGCTAA GAGAAACTCT CAGAGAAATG AAGGGGTTCC ACCAGGCCAT GGGCAGCCAC TAGAGCCAAC CCTTGGAGGA GTTTGACTCC ACTGAGCCTT GGTGTGGTGT	22146 22206 22266 22326 22386 22446 22506
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG Gly Ser Gly Thr GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC ACTCAGCTAA GAGAAACTCT CAGAGAAATG AAGGGGTTCC ACCAGGCCAT GGGCAGCCAC TAGAGCCAAC CCTTGGAGGA GTTTGACTCC ACTGAGCCTT GGTGTGGTGT	22146 22206 22266 22326 22386 22446 22506 22566
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC ACTCAGCTAA GAGAAACTCT CAGAGAAATG AAGGGGTTCC ACCAGGCCAT GGGCAGCCAC TAGAGCCAAC CCTTGGAGGA GTTTGACTCC ACTGAGCCTT GGTGTGGTT TTCCATCTGT GAGATGGGAA TACTTTGCCC AAGAGCCTGT TAGAAGCTGT AGGAAGCACA GAGTCGGCTA GGTATAGATT TGCTCTCACC TCCATCTCTC GATACCAGTT CTCTGCAGAG CTTGGGTTTG TGGGAGGGGT GGGGGGGTGA GGGGAGAAGG CTGTGAGCTG CAGCTAGCCA GAGGGGTCTC	22146 22206 22266 22326 22386 22446 22506 22566
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 40 45 GGC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC ACTCAGCTAA GAGAAACTCT CAGAGAAATG AAGGGGTTCC ACCAGGCCAT GGGCAGCCAC TAGAGCCAAC CCTTGGAGGA GTTTGACTCC ACTGAGCCTT GGTGTGGTT TTCCATCTGT GAGATGGGAA TACTTTGCCC AAGAGCCTGT TAGAAGCTGT AGGAAGCACA GAGTCGGCTA GGTATAGATT TGCTCTCACC TCCATCTCTC GATACCAGTT CTCTGCAGAG CTTGGGTTTG TGGGAGGGGT GGGGGGGTGA GGGGAGAAGG CTGTGAGCTG CAGCTAGCCA GAGGGGTCTC CCAGAAGAAT GGGGAGAGCT AAGAAGGAAA GTTGAGGTCA CAGTGGGAAG GAGACCAGAG	22146 22206 22266 22326 22386 22446 22506 22566 22626
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser Gly Ser Gly Thr GCTGGAGTGG TGAGCAGGCA GTCACCCAGC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC ACTCAGCTAA GAGAAACTCT CAGAGAAATG AAGGGGTTCC ACCAGGCCAT GGGCAGCCAC TAGAGCCAAC CCTTGGAGGA GTTTGACTCC ACTGAGCCTT GGTGTGGTT TTCCATCTGT GAGATGGGAA TACTTTGCCC AAGAGCCTGT TAGAAGCTGT AGGAAGCACA GAGTCGGCTA GGGCAGGCGTA GGGGAGGGT GGGGGGGTGA GGGGAGAAGG CTGTGAGCTG CTCTGCAGAG CTTGGGTTTG TGGGAGGGGT GGGGGGGTGA GGGGAGAAAG CTTGAGGTCA CAGTGAGCCA GAGGGGTCTC CCAGAAGAAT GGGGAGAGCT AAGAAGGAAA GTTGAGGTCA CAGTGGGAAG GAGACCAGAG CAAAAGGGTT GAAGGGTTG GAAGGTTG GAAGGTTG GAAGGTTG TTGGGAGCCTT AGGACCAGAG CAAAAGGGTT GAAGGTAGGT AAAATGCAGC CGTGTATTCT TGGGAGCCTT AGGCCTTGGG	22146 22206 22266 22326 22386 22446 22506 22566 22626 22686
Pro Pro Glu Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser 30 GCC TCT GGC ACA GGTAAGACTG ACCCAGAACA CTGAGATGGC ATAGATCATG GLY Ser Gly Thr GCCCAGCC TTTTAGTGAA CCCCCTTCTT CTCCCATCCC ATCCTTAGCC ATTGGAGTCA GGACAGTGCC AAAAGGAAGA ATGGTATCCA GCTGCAAGCC ACCCAGCTAA GAGAAACTCT CAGAGAAATG AAGGGGTTCC ACCAGGCCAT GGGCAGCCAC TAGAAGCCAAC CCTTGGAGGA GTTTGACTCC ACTGAGCCTT GGTGTGGTT TTCCATCTGT GAGATGGGAA TACTTTGCCC AAGAGCCTGT TAGAAGCTGT AGGAAGCACA GAGTCGGCTA GGGAGGGGT GGGGGGGTGA GGGGAGAAGG CTGTGAGCTG CAGCTAGCCA GAGGGGTTTC CAGAAGAAT GGGGAGAGACT AAGAAGGAAA GTTGAGGTCA CAGTGGGAAG GAGACCAGAG CAAAAGGGTTG GAAGGTAGGT AAAAATGCAGC CGTGTATTCT TGGGAGCCTT AGGCCTTGGG CAAGAGGGTA GAAGAGGGTA TTGTCCTGGG CTGCAGTCCT TGGGAGCCTT AGGACCAGAG CAAAAGGGTAG GAAGAGGAAA GTTGAGGTCA GAGTCGCTT AGGCCTTGGG CAAAAGGGTAG GAAGAGGAAA GTTGAGGTCA GAGTCAGCT AGGCCTTGGG CAAGAGGGTA GAAGAGGATA TTGTCCTGGG CTGCAGTCCT TGGGAGCCTT AGGCCTTGGG CAAGAGGGTA GAAGAGGATA GTTGAGGTCA GAGTCAGCT TGGGTTTTCT TGGGAGCCTT AGGCCTTGGG CAAGAGGGTA GAAGAGGATA GTTGAGGTCA GAGTCAGCT TGGGTTTTCT TGGGAGCCTT AGGCCTTGGG CAAGAGGGTA GAAGAGGGTA GAAGAGGAAA GTTGAGGTCC TGGTGTCTTGG CAAGAGGGTA GAAGAGGGTA GAAGAGGAAA GTTGAGGTCC TTGGGAGCCTT AGGCCTTGGG CAAGAGGGTA GAAGAGGATA GTTGCTCTGGG CTGCAGTCCT TGGGAGCCTT AGGCCTTGGG CAAGAGGGTA GAAGAGGATA GTTGAGGTCC TTGGGAGCCTT AGGCCTTGGG CAAGAGGATA GAAGAGGATA GTTGAGGTCC TTGGGAGCCTT AGGCCTTGGG CAAGAGGATA GAAGAGGATA GTTGAGGTCC TTGGGAGCCTT AGGCCTTGGG CAAGAGGATA GAAGAGGATA GTTGAGGTC TTGGGAGCCTT AGGCCTTGGG CTGCAGAGAGAGA GAAGAGGATA GTTGAGGTC TTGGGAGCCTT AGGCCTTGGG CTGAGAGAGAA GAAGAGGAAA GTTGAGGTC TTGAGGTCA GAGAGCACA GAGAGAGAA GTTGAGGTA GAAGAGAA GTTGAGGTA GAAGAGAA GTTGAGGTA GAAGAGAA GTTGAGGTA GAAGAGAA GTTGAGAGAA GTTGAGAGAA GTTGAGAGAA GTTGAGAGAAA GTTGAGAGAA GTTGAGAGAAA GTTGAGAGAAA GTTGAGAGAAA GTTAAAAA GAAAGAAA	22146 22206 22266 22326 22386 22446 22506 22566 22626 22686 22746



TGAGTTCAGC CTTTGCTAGG TCACCTTTGG GGTCTCAGAA GGCTTCAGCT CCTGGTAGAG	22926
CATGAATCAC GTCAGGCGTG ATGCTGGAGA CCTCTCCTAC CCTGACACCC CAAACCCCCA	22986
CCTCTGACCC TGCA GGT GCT TTG CCA GAT ACT TTG TCA CGG CAG ACA CCT Gly Ala Leu Pro Asp Thr Leu Ser Arg Gln Thr Pro 50 55 60	23036
TCC ACT TGG AAG GAC GTG TGG CTG TTG ACA GCC ACG CCC ACA GCT CCA Ser Thr Trp Lys Asp Val Trp Leu Leu Thr Ala Thr Pro Thr Ala Pro 65 70 75	. 23084
GAG CCC ACC AGC AGC AAC ACC GAG ACT GCT TTT ACC TCT GTC CCA Glu Pro Thr Ser Ser Asn Thr Glu Thr Ala Phe Thr Ser Val Leu Pro 80 85 90	23132
GCC GGA GAG AAG CCC GAG GAG GGA GAG CCT GTG CTC CAT GTA GAA GCA Ala Gly Glu Lys Pro Glu Glu Gly Glu Pro Val Leu His Val Glu Ala 95 100 105	23180
GAG CCT GGC TTC ACT GCT CGG GAC AAG GAA AAG GAG GTC ACC ACC AGG Glu Pro Gly Phe Thr Ala Arg Asp Lys Glu Lys Glu Val Thr Thr Arg 110 125	23228
CCC AGG GAG ACC GTG CAG CTC CCC ATC ACC CAA CGG GCC TCA ACA GTC Pro Arg Glu Thr Val Gln Leu Pro Ile Thr Gln Arg Ala Ser Thr Val 130 135 140	23276
AGA GTC ACC ACA GCC CAG GCA GCT GTC ACA TCT CAT CCG CAC GGG GGC Arg Val Thr Thr Ala Gln Ala Ala Val Thr Ser His Pro His Gly Gly 145 150 155	23324
ATG CAA CCT GGC CTC CAT GAG ACC TCG GCT CCC ACA GCA CCT GGT CAA Met Gln Pro Gly Leu His Glu Thr Ser Ala Pro Thr Ala Pro Gly Gln 160 165 170	23372
CCT GAC CAT CAG CCT CCA CGT GTG GAG GGT GGC GGC ACT TCT GTC ATC Pro Asp His Gln Pro Pro Arg Val Glu Gly Gly Gly Thr Ser Val Ile 175 180 185	23420
AAA GAG GTT GTC GAG GAT GGA ACT GCC AAT CAG CTT CCC GCA GGA GAG Lys Glu Val Val Glu Asp Gly Thr Ala Asn Gln Leu Pro Ala Gly Glu 190 195 200 205	23468
GGC TCT GGA GAA CAA GTGAGTGGCT TTGCATTTCC TGGGTGGCCA CTAGTGCCTG Gly Ser Gly Glu Gln 210	23523
CACCTGGCCG CCTAATGTCC TCATTACAGT GACAGGTGAC AGGGTCCCAC CTTCCTCCTG	23583
CCCGAAACAG ACTGATTGCA AGATCAGGAG GTGGGCGACT CCTTAGATGT CATTCAGGAG	23643
CTTACAGCAG GGTGAATTTT CCGTCTTAGA CCTTCATGGG AATTTTCACA CAACAATGTG	23703
TACGTTGTGT CACTGGAGGC GGTATCTGTG TCTTGGCCTG CCAGGGTCCC AGGTGTGACT	23763
GACTGCATTC CTTGACAGAT GCTGGTATAG GTTGGCTACG TCTGATGGGG GTGGCAGGGG	23823

ATCCCATCAG GTATGGCACT GCTCAGGTTG CTGTTGTGTC AGTGGCTCCA GCTGACCTGA	23883
TCCCAACCTA CCCTTCTGTA G GAC TTC ACC TTT GAA ACA TCT GGG GAG AAC Asp Phe Thr Phe Glu Thr Ser Gly Glu Asn 215 220	23934
ACA GCT GTG GCT GCC GTA GAG CCC GGC CTG CGG AAT CAG CCC CCG GTG Thr Ala Val Ala Ala Val Glu Pro Gly Leu Arg Asn Gln Pro Pro Val 225 230 235	23982
GAC GAA GGA GCC ACA GGT GCT TCT CAG AGC CTT TTG GAC AGG AAG GAA Asp Glu Gly Ala Thr Gly Ala Ser Gln Ser Leu Leu Asp Arg Lys Glu 240 245 250	24030
GTG CTG GGA GGTGAGTCTT CTTTCAGGTG GAGAGGAGGA GGCAGGTGGT Val Leu Gly 255	24079
GGCTCTGAGG TAGCCTGGGT TGCTGGGGTG AAGCATCTTT AGCAGCAGGG TGGGGAAGGA	24139
GGAGGGTCAA TTCTACTCCA GGCCACCTCC TAGGCTGTCC GTCTAGTCTG GGAGAGACTA	24199
CCACTGACCC CGTGGAGCTA CTGATCTGAG CCTGCCTCTG TTCACTCCCT A GGT GTC Gly Val	24256
ATT GCC GGA GGC CTA GTG GGC CTC ATC TTT GCT GTG TGC CTG GTG GCT Ile Ala Gly Gly Leu Val Gly Leu Ile Phe Ala Val Cys Leu Val Ala 260 265 270	24304
TTC ATG CTG TAC CGG ATG AAG AAG GAC GAA GGC AGC TAC TCC TTG Phe Met Leu Tyr Arg Met Lys Lys Asp Glu Gly Ser Tyr Ser Leu 275 280 285	24352
GAG GAG CCC AAA CAA GCC AAT GGC GGT GCC TAC CAG AAA CCC ACC AAG Glu Glu Pro Lys Gln Ala Asn Gly Gly Ala Tyr Gln Lys Pro Thr Lys 290 295 300 305	24400
CAG GAG GAG TTC TAC GCC TGATGGGGAA ATAGTTCTTT CTCCCCCCAC Gln Glu Glu Phe Tyr Ala 310	24448
AGCCCCTGCC ACTCACTAGG CTCCCACTTG CCTCTTCTGT GAAAAACTTC AAGCCCTGGC	24508
CTCCCCACCA CTGGGTCATG TCCTCTGCAC. CCAGGCCCTT CCAGCTGTTC CTGCCCGAGC	24568
GGTCCCAGGG TGTGCTGGGA ACTGATTCCC CTCCTTTGAC TTCTGCCTAG AAGCTTGGGT	24628
GCAAAGGGTT TCTTGCATCT GATCTTTCTA CCACAACCAC ACCTGTTGTC CACTCTTCTG	24688
ACTTGGTTTC TCCAAATGGG AGGAGACCCA GCTCTGGACA GAAAGGGGAC CCGACTCTTT	24748
GGACCTAGAT GGCCTATTGC GGCTGGAGGA TCCTGAGGAC AGGAGAGGGG CTTCGGCTGA	24808
CCAGCCATAG CACTTACCCA TAGAGACCGC TAGGTTGGCC GTGCTGTGGT GGGGGATGGA	24868
GGCCTGAGCT CCTTGGAATC CACTTTCAT TGTGGGGAGG TCTACTTTAG ACAACTTGGT	24928

WO 94/12162



TTTGCACATA TTTTCTCTAA TTTCTCTGTT CAGAGCCCCA GCAGACCTTA TTACTGGGGT 24988 AAGGCAAGTC TGTTGACTGG TGTCCCTCAC CTCGCTTCCC TAATCTACAT TCAGGAGACC 25048 GAATCGGGGG TTAATAAGAC TTTTTTTGTT TTTTGTTTTT GTTTTTAACC TAGAAGAACC 25108 AAATCTGGAC GGCAAAACGT AGGCTTAGTT TGTGTGTTGT CTCTGAGTTT GTCGCTCATG 25168 CGTACAACAG GGTATGGACT ATCTGTATGG TGCCCCATTT TTGGCGGCCC GTAAGTAGGC 25228 TGGCTAGTCC AGGATACTGT GGAATAGCCA CCTCTTGACC AGTCATGCCT GTGTGCATGG 25288 ACTCAGGGCC ACGGCCTTGG CCTGGGCCAC CGTGACATTG GAAGAGCCTG TGTGAGAACT 25348 TACTCGAAGT TCACAGTCTA GGAGTGGAGG GGAGGAGACT GTAGAGTTTT GGGGGAGGGG 25408 TGGCAAGGGT GCCCAAGCGT CTCCCACCTT TGGTACCATC TCTAGTCATC CTTCCTCCCG 25468 GAAGTTGACA AGACACATCT TGAGTATGGC TGGCACTGGT TCCTCCATCA AGAACCAAGT 25528 TCACCTTCAG CTCCTGTGGC CCCGCCCCCA GGCTGGAGTC AGAAATGTTT CCCAAAGAGT 25588 GAGTCTTTTG CTTTTGGCAA AACGCTACTT AATCCAATGG GTTCTGTACA GTAGATTTTG 25648 CAGATGTAAT AAACTTTAAT ATAAAGGAGT CCTATGAACT CTACTGCTTC TGCTTCTTCT 25708 TCTCTGGACT GGTGGTATAG ATATAGCCAC CCTTTGCCCA AACCCTGGTA GCTCGGGGAA 25768 GCTTGGCTTA AGGCTGCACG CCTCCAATCC CCCAAAGGTA GGATCCTGGC TGGGTCCAGG 25828 25888 TTTTGGAAGT TGGTAAGTTC AGCCAAGGTT TTACAGGCCC TGATGTCTGT TCTTCTAAAT 25948 GGTTTAAGTA ATTGGGACTC TAGCACATCT TGACCTAGGG TCACTAGAGC TAAGCTTGCT 26008 TTGCAGGGCA GACACCTGGG ACAGCCTTCC TCCCTCATGT TTGCTGGGAC ACTGCTGAGC 26068 ACCCCTTGCT TACTTAGCTC AGTGATGTTC CAGCTCCTGG CTAGGCTGCT CAGCCACTCA 26128 GCTAGACAAA AGATCTGTGC CCTGTGTTTC ATCCCAGAGC TTGTTGCCAG ATCACATGGC 26188 TGGATGTGAT GTGGGGTGGG GGTGGGGTCA TATCTGAGAC AGCCCTCAGC TGAGGGCTTG 26248 TGGGACAGTG TCAAGCCTCA GGCTGGCGCT CATTCATATA ATTGCAATAA ATGGTACGTG 26308 TCCATTTGGA CAGCAGACAC TTTGGTGTAC TTGTGCAGTC TCTTTTTGGT CTGGACCATG 26368 TCCAACTCTA TCTGGTTTTT GGAATGGGAG CCTAACTGGC CTGTGTTCTG GCTTGGTACC 26428 AAATAGCAAC AGTCAGTGGC ATCCTTGCCC AGGCCCCAGG GCAGGACTAT GCTCTTGCCA 26488 TATCCAGGAC TCCCGACTTT GCACCTGTTT TCCCTCTGTG TGTAGCATCA TGAACTCCAG 26548 CTAGGTTGTT CCTTTCCCTG GGGTCAGGAG GATTCTGCTG ACTCTGAATG TCAGGATTTG 26608 CTTTTGTTCT GTTTGCTTAT TGGGCAATTC TCAACCTTCA CTAGCAACAG TCTCATGTGT 26668

CAGGATTACA AGTATTGCTT GCACATTGAG GG

26700

(2) INFORMATION FOR SEQ ID NO: 2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 311 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (x1) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Met Arg Arg Ala Ala Leu Trp Leu Trp Leu Cys Ala Leu Ala Leu Arg
1 5 10 15

Leu Gln Pro Ala Leu Pro Gln Ile Val Ala Val Asn Val Pro Pro Glu 20 25 30

Asp Gln Asp Gly Ser Gly Asp Asp Ser Asp Asn Phe Ser Gly Ser Gly 35 40 45

Thr Gly Ala Leu Pro Asp Thr Leu Ser Arg Gln Thr Pro Ser Thr Trp 50 55 60

Lys Asp Val Trp Leu Leu Thr Ala Thr Pro Thr Ala Pro Glu Pro Thr 65 70 75 80

Ser Ser Asn Thr Glu Thr Ala Phe Thr Ser Val Leu Pro Ala Gly Glu 85 90 95

Lys Pro Glu Glu Glu Pro Val Leu His Val Glu Ala Glu Pro Gly 100 105 110

Phe Thr Ala Arg Asp Lys Glu Lys Glu Val Thr Thr Arg Pro Arg Glu 115 120 125

Thr Val Gln Leu Pro Ile Thr Gln Arg Ala Ser Thr Val Arg Val Thr 130 135 140

Thr Ala Gln Ala Ala Val Thr Ser His Pro His Gly Gly Met Gln Pro 145 150 155 160

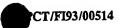
Gly Leu His Glu Thr Ser Ala Pro Thr Ala Pro Gly Gln Pro Asp His 165 170 175

Gln Pro Pro Arg Val Glu Gly Gly Gly Thr Ser Val Ile Lys Glu Val
180 185 190

Val Glu Asp Gly Thr Ala Asn Gln Leu Pro Ala Gly Glu Gly Ser Gly 195 200 205

Glu Gln Asp Phe Thr Phe Glu Thr Ser Gly Glu Asn Thr Ala Val Ala 210 215 220

Ala Val Glu Pro Gly Leu Arg Asn Gln Pro Pro Val Asp Glu Gly Ala 225 230 235 240



Thr Gly Ala Ser Gln Ser Leu Leu Asp Arg Lys Glu Val Leu Gly Gly 245 250 255

Val Ile Ala Gly Gly Leu Val Gly Leu Ile Phe Ala Val Cys Leu Val 260 265 270

Ala Phe Met Leu Tyr Arg Met Lys Lys Lys Asp Glu Gly Ser Tyr Ser 275 280 285

Leu Glu Glu Pro Lys Gln Ala Asn Gly Gly Ala Tyr Gln Lys Pro Thr 290 295 300

Lys Gln Glu Glu Phe Tyr Ala 305 310

(2) INFORMATION FOR SEQ ID NO: 3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2196 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: both
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iii) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

TCTAGAACAC TTATTAAGAG CCAGGCACTG AAAAGTGCAG ACTCCCTCAT TTCATCCTGG 60 CCGTGCTTAC AAGTAGTTTC CATGCTCTGG TAACCCTGTG CAGAGGGCAG CGTGGGAGGC 120 GGGCCGCTTG GTGGACGGTC ATGGGGGCTC TGCATGGGTG GTTGCCCTTG CCTCAGAAGA 180 ACTCCCTAAG TAAGAGCAAG TTAGCCTCCC TAACCCCTGG TGGGTTGTTG CTTCTTTCT 240 300 CCTCTTGTTT CTGCCAAGAG AGGGTGGACC AAGAAGACCC CAGCCTACAG AACATGTGAT CCAAATAAAC TTCTTTTTAG TATAAATGTC CTAGCCTGTG ACGTTCTGGT AGACTAGCAC 360 AAGATGGACC AAGACAACTC TCATCGAGAC TCTGAGGAAC GAACTGGCAT CACATGGGAA 420 CAGGAAATGA AGCTTAGAGA GAGGTTCTGT GGCTTGTCCA ACATGGCTGT AGTTTAAATC 480 540 GGGCTCCGGA GCTAGAAGGA CACGTGTATC AGCCATGGCT TCAGTTTATT GCTGTATACT 600 CTGTGCTTCT GGCTCTCATG GAAAAGACAG ACATTGGGGT TCTTATAATC TCTCCCTCTC 660 CCCTCCCCAC ACTCTATCCC CAAAGGAGGC ACCACTTCTG CAGGTAAATG TTATCTTCAA 720 AGCGCTCACA TCGCAACCTT TGCCCACACC ATCTCATTAA AGGAATTGGC AGTGACTTTA 780

AGGTGAAAGA	ACTCGGTGGC	TACGTGTTAT	ATAAATTTGC	ATCTGGGTCT	CAGAGCTGGA	840
AGGAAGGCAC	TCCCATACAT	GCAGTCTGTA	CATGCAGTCG	GATGATGGAC	CAACAACACA	900
TTGTGATTTA	TGCCCCTGCT	GGTGAGCCCA	GGAATCCCTG	TAGCACTCTC	TCTCAGCTCT	960
AGGGCCCTGC	TTGTGTATGG	AAAACGCTTA	GTGTTTTATA	GGTATTTTGT	CAGAATACTT	1020
TAAGGAACTT	GACCAAAGTT	ACAGGGAGGT	TAGACAGATT	GTCATGGTAT	ACTCACCTCT	1080
GTCTCTGACC	CTCCTAACTG	GGACCTCTTT	AGTÇTCCCTT	GAGGCAGGGA	GTGCCACATG	1140
CATGTGTCCA	GGCACATGTC	TCCTGGTTTA	CCTCCCAACG	CACCTCAAGT	CCCCAAGGTA	1200
GGTAGGCACT	TGTATTCTGT	AATTCAGAGA	GGCAAATCAA	ACTGTTACAA	TGTTTGCCCA	1260
AAGCTCCCCA	AGCAAAGTGG	CCCTAAGAGT	GAGCAAAGAG	ACTGCGTGCC	TTCACTGCCT	1320
GTGTGAATCC	CTGCAGATAG	TCTCTCATCT	TGGTGCCCTT	CCCACAGAGG	CTGGGGCGGC	1380
AGGAGGGAGC	CTGGACAGCT	CAGACACTGG	GTCATTGATG	ACTGTTGTGT	GGGATACCTG	1440
CCGGGGCGCA	GGAGTGAGCC	ATGCCACCCC	AGGAAGTGGT	TCAGGGTGAC	TCTTCTTGGC	1500
ACACCTGGGA	GGATGTAGCT	GGTGCTGGCA	CACCCACCGT	CACGAGAGCT	TCCTGTCCAA	1560
ACCTTCAACA	AAGGCGGCTT	CTTGAGACAG	GCTAGACTGA	AGTCACCAGC	CTTGGGTGGG	1620
GTCCACTATG	TAACCTCAGT	GCTCAGGAAC	CCTTTCCCAT	ACTGTCTGGA	ACTATACTGT	1680
ATGTAGCTGG	GTTTCCACGC	ATGTGTGCCT	GCACCCAGTC	CATCTCATCT	TCTATCTCCC	1740
TCCCCTTTCC	CGCTTCCCCC	CTCCCCACTC	TCCATCTCAT	CTTCCATCCC	CACCTCTTCT	1800
GGTCCCTGCC	CTGCTAAACT	CAGGGTAGCT	GCATTCCGCT	GGCCTTCCCC	ATGTTCCAGG	1860
CTTCAGTCCC	TTCTCTGCAC	CTGTCCTTTG	TGAAGTGACC	AGAGGATTTC	TGATCCTGTC	1920
TCTGTCGCTC	TGAAGGGTCA	GGAGTTCCTC	CTGCCTGGAC	AAAGCCATCC	TGACGCACAT	1980
АААТААААСА	AACATCAAAC	TCTATTCAAC	CCCCTGGAAC	CCGTGTGTGT	TACTTACAGG	2040
GCAAAAGAAT	GGAGCAGGGG	ATGGGTTGTG	GGGGGGGGG	GTGGCATCTG	GGTTGTCTAC	2100
AGTTGTGCAT	TAAGTTGTAA	TTAAGATGTG	CATTTCTCCA	AATAAGGGAA	AATTATTCTG	2160
GATTATTTGA	GTGAAGCTGA	AAGGTGATCA	TCTAGA			2196

WHAT IS CLAIMED IS:

- 1. A method of decreasing the growth of a malignant cell wherein said method comprises exposing said cell to efficacious levels of a compound that induces expression of syndecan in said cell.
- 2. The method of claim 1, wherein said induced expression is obtained by affecting the enhancer element of the syndecan gene (SEQ ID No. 3).
- 3. The method of claim 1, wherein said induced expression is obtained by affecting the suppressor element of the syndecan gene.
 - 4. The method of claim 1,2 or 3, wherein said cell is steroid-responsive.
 - 5. The method of claim 4, wherein said steroid is estrogen or androgen.
- 6. The method of claim 4, wherein said cell is selected from the group consisting of a malignant breast cell, an endometrium cell and a prostate cell.
- 7. The method of claim 6, wherein said method further comprises exposing said cell to an anti-steroid agent.
 - 8. The method of claim 7, wherein said agent is toremifene or tamoxifen.
 - 9. The method of claim 1,2 or 3, wherein said cell is a mesenchymal cell.
 - 10. The method of claim 6 or claim 9, wherein said cell is a human cell.
- 11. The method of claim 1, wherein said cell is exposed to efficacious levels of a composition comprising growth factors of said cell.
- 12. The method of claim 11, wherein said growth factors are selected from the group consisting of bFGF; TGF-ß; and bFGF and TGF-ß.
- 13. A method for treating a patient in need of treatment to decrease the growth of a malignant tumor in said patient, wherein said method comprises administering, to said patient, efficacious levels of a composition that induces expression of syndecan in the cells of said tumor.



- 15. A method of claim 13, wherein said composition affects the suppressor element of the syndecan gene.
- 16. The method of claim 13, 14 or 15, wherein said tumor is a tumor whose growth is stimulated by steroids.
 - 17. The method of claim 16, wherein said steroid is estrogen or androgen.
- 18. The method of claim 17, wherein said tumor is selected from a tumor the group consisting of a breast, endometrium, prostate gland or mesenchymal tissue.
- 19. The method of claim 13, wherein said composition comprises an antisteroid agent.
 - 20. The method of claim 19, wherein said agent is toremifene or tamoxifen.
- 21. The method of claim 13, wherein the efficacious agents in the composition comprise growth factors.
- 22. The method of claim 21, wherein said growth factor is selected from the group consisting of bFGF; TGF-B; and bFGF and TGF-B.
 - 23. The method of claim 13, wherein said patient is a human.
- 24. A method of inducing differentiation of a cell to a more differentiated phenotype, wherein said method comprises exposing said cell to efficacious levels of a composition that induces expression of syndecan in said cell.
- 25. The method of claim 24, wherein the composition affects the enhancer element of the syndecan gene.
- 26. The method of claim 24, wherein the composition affects the suppressor element of the syndecan gene.

- 27. The method of claim 24, wherein the growth of said cell is stimulated by steroids.
 - 28. The method of claim 27, wherein said steroid is estrogen or androgen.
 - 29. The method of claim 24, wherein said cell is an epithelium cell.
- 30. The method of claim 29, wherein said epithelium cell is an epidermal skin cell.
 - 31. The method of claim 24, wherein said cell is a human cell.
- 32. The method of claim 24, wherein said composition comprises an antisteroid agent.
- 33. The method of claim 24, wherein the efficacious agent in said composition is toremifene or tamoxifen.
- 34. The method of claim 24, wherein the efficacious agent in said composition is a growth factor.
- 35. The method of claim 34, wherein said growth factor is selected from the group consisting of bFGF; TGF-B; and bFGF and TGF-B.
- 36. A method of stimulating hair growth in an epidermal cell of skin, wherein said method comprises stimulating expression of syndecan in said cell.
- 37. The method of claim 36, wherein sald stimulated expression is achieved by affecting the enhancer element of the syndecan gene.
- 38. The method of claim 36, wherein said stimulated expression is achieved by affecting the suppressor element of the syndecan gene.
 - 39. The method of claim 36, wherein said cell is a mesenchymal cell.
- 40. The method of claim 36, wherein said method comprises administering a composition comprising efficacious levels of a growth factor.

PCT/FI93/00514

41. The method of claim 40, wherein said growth factor is selected from the group consisting of bFGF; TGF-B; and bFGF and TGF-B.

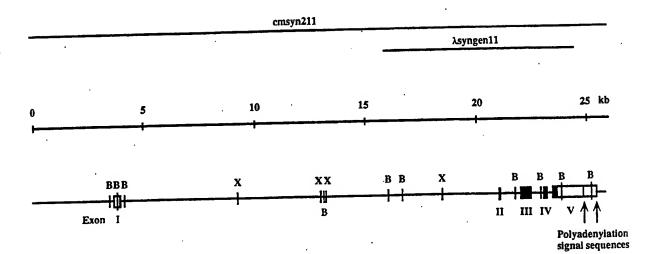
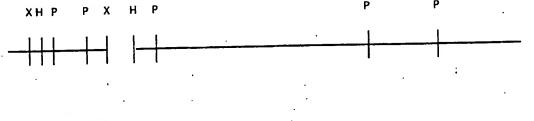
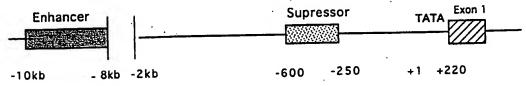


Fig 1.





X= Xba I H= Hind III P= Pst I

Fig 2.

Fia 3.

-4078 gtctttcctctgtcttctga ctcagatgcttagctagctc tttaggacccaccctcacac ctgcaaataatactttattt -3918 aaaagtaaaataaattaaaa aatagaaaggtttgagcatg atggcccagtggtaaaggcc agtggctccaacgcaagtcc -3838 tgacaaatggtaacgggcct gttcttcaggcttgagggaa gtttattgattgaggctaaa agcaacccaaaggctccact -3758 tgcctagtgtgaagccctgg atgtgctctcccacactgca tgtccacctgtggtgtcagc acctgggaagctgaggatga -3598 attgtaatcccagcacttga cagaccaatggggggggat tgctgtgagtttaagacagc ctggcctacaaagaaaaacc -3518 ctacccaaacccaagaaaaa tgaaaccagtaatataaata gctattttcattttaaatgc tctaaagacacagcgttaac -3438 acaaaagctctcgtctgtgg ttcctattccctccttctcc cccaggtcttctttaatgta tactttttgtttgcttattt -3358 gcttgttttggattttggct tttaaagacagggtctcact atgtagctccaactatttgg gaactcactatgtagaccag -3278 gctagccagggacttataga gatctacctaccactgcctc ccaagtgctgagactaaagg catgtgacactttgcttggt -3198 tattacaaacattttaaaag aacattttgaacattaatag atgtatgtatatatatcact ctatgtagtatatatgttag -3038 tttttttattatttatta ttatatgtaagtacactgta gctgtcttcagacanaccag aagagggagtcagatcttgt -2958 tacggatggttgtgagcacc atgtggttgctgggattcga actctggaccttccgaagag cagtcgggtgctcttaccca -2878 ctgagccatctcaccagccc cttaaatttattttatctt atgtccattggtgttttgcc tgcatgtatgtgtaaaagtg -2798 tcagaaactgaagttacaga ctgttgtgagctaccattgt tgtgggtgctgggacttgaa cctgggtcctctggaagagc -2718 agreattattettaaceact gagecatetetetagecete gttttttagttttttttt gttttgttttttttt -2638 titttaagattttcttattt attatatgtaagtacactgt agctgtcttcagacactcca gaagagggcgccagatctcg -2558 ttatggatggttgtgagcac catgtggttgctgggaattg aactccagacctttggaaga gcagtcagtgctcttaactg -2478 ctgagccatctctccagccc cgttttttaggtttttgaag acagggtttcctgtgtagct ctagctgtccaggaactagc -2398 tetgtagaccaggttggeet caaatttagagatttgeetg tetetetgeetetegagage tgggattaaaagtgtgeage -2318 ccaacaatctactcaaagta ggttttgaaaaagctttcca tattaggagttaactagctt catttcagaaatactgcatg -2238 gaattcaaatgtgggaccat tcatagctactttggttttc cttcagtgacaggcattcgg catgcctattagggaagtca -2158 aatggcctggagaagtcatc ctgggtgagagggctaatgc attttcagcttgacagacac tgtcaacctatgcagacagt -2078 ctgctccagctcagatgtca attgcatgcagacctgcagt cagacgctaagctccctacc tactctccatcagcttagat -1918 tettattttatgtaaattgg tacttcacttacatgtatgt ccgtgtgaggatgttgtatc ctctggtactggagttatag -1838 acagetgtaagtegecatac aggtgetgggaattgaaece tgateetetggaagaatagt cagtgetettaaeceetgag -1758 ccatctctccaacctcttgc atattgaggacagggaggaa tcacaagccatgtagggtgc ctgggctctgaggtcaacag -1678 gaccatagcctcctttcttt atgtgcctttcttggggtct ccctataggagtcgtcttcg ttgcctctttactgtctcat -1598 tgatctgggctaaacttatg cagttggaaggaaagatcaa gctggtcatgtttaaaacat gaaacagcctcatcagttcc -1518 cttcctgttcccgtctcccc ccccctcccgccccattt tgagaggacaggaaggtaaa ataccaaagtgtcctatttt -1438 cctccaaatatcaggctcaa aggactgaagagctgacttc agatcccaaagccactgtgt taggaggcacctgctttta -1358 ggtcctaagccttcctgagc cttgctattgggtattcttt accaagaccctcaaggatct aggcaagaactgggcaggat -1278 ctgtatgtagcccatagtta gacctagggcagctgagacg ccaaaagggagagtttcctg aggacaaaagtgttcaaaca -1198 caactgggtgctggttgttg ggctactcgtggaggtgtgg tgtgtgtaaaggaggctgtt gaattcccagaaggctggtt -1118 ccacagtgtagagtctacac tggggacttcccgagacgct gagcctcagatctagcttct cagtccaggccagctgatgt -1038 ggggctgaggaacaaggatg gatgccatctatggccctgc cttgcaggtgcaaagggcct ttggcaccatctacagattg -958 agggcaagacagggctggtt cttcctccttgctctcgctg ctatctgcctcgcctgtagg ctctctgggctcctttttgg -878 actgacacgtctgaaggagc ttggaaactgtgaggtccag gccccatagagaatcatgaa ggaacaggaattcaactgga -798 geteegeagetggttaggee tgeggtcacetggaaacaaa gaggccatttattttteet ttggtcttggacaaggaaga -718 gaaggggctttctataaata gaaagacagcaaaaaagaaa ataataataataataat aataataataataataaaa -638 caataacaaagccagctctt ccagacagtgctcatgtctt taaaggtctttaaaggtctg gagttcccagcaattaagta -558 aaggaccaagacctcagggg tecectatectcagecegtg gggaggtgggaaccatacat egateceteggtttatatat -478 agecteategetgtgggget eegaggttgeeeccaaaate ttgeteacetggaggaeece tgggtgteetegeecagagg -398 gegetgeagectegeacgta gagaactaacategeeette tecagggeagtgeeteegga eteeggaecaggaeatagta -318 gcgagtgcacctgggtctcc gtcagctacgcatcaaggaa ggtgcgacgcggggaattaca gattgccggcactcaccagt -158 ggtgtggctggatccctggg gggtggggcgctccaagggg cggggcaacccagggggcgg ggcccgaggggtggagattg -78 ggactacccaggcccgcgga gctgggggtgggcggctagt tttgcaactgcagagccttt gggtttattataaggcggAG

tctagatattcaaactcac cagatggagtgatgtccacc cctattggtgggagtgacta 3 CTCCGCGGGAGAGGTGCGGG CCAGAGGAGACAGAGCCTAA CGCAGAGGAAGGGACCTGGC AGTCGGGAGCTGACTCCAGC 83 CGGCGAAACCTACAGCCCTC GCTCGAGAGAGCAGCAGCT GGGCAGGAGCCTGGGACAGC AAAGCGCAGAGCAATCAGCA

163 GAGCCGGCCCGGAGCTCCGT GCAACCGGCAACTCGGATCC ACGAAGCCCACCGAGCTCCC GCCGCCGGTCTGGGCAGCAT



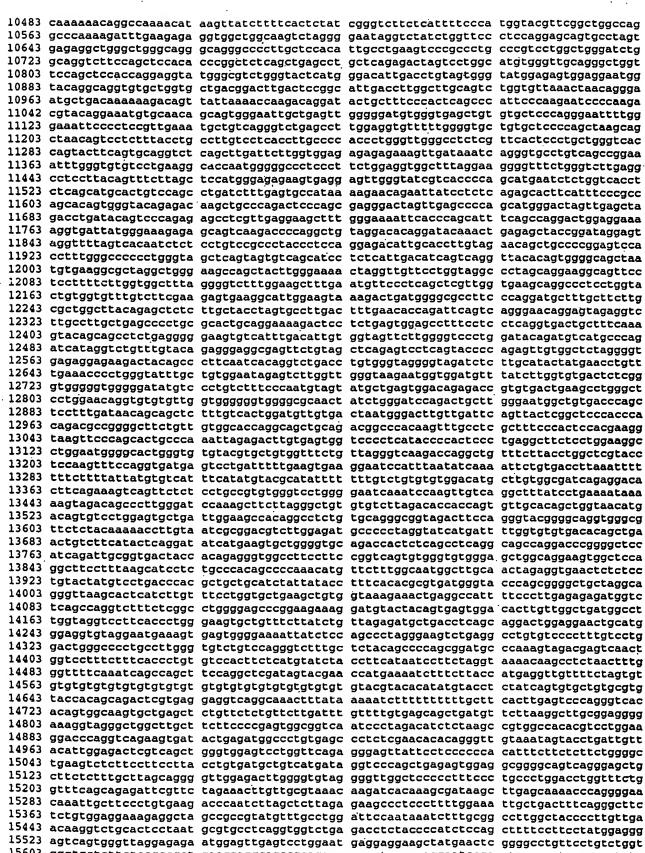
243 GAGACGCGCGCGCTCTGGC TCTGGCTCTGCGCGCTGGGG CTGCGCCTGCAGCCTGCCCT CCCGgtgagtgtggcccggg targargalaalaLeuTrpL euTrpLeuCysalaLeuAla LeuargLeuGlnProAlaLe uPro

323 gcagggctgggaggcggcgg gaagccgggactcgccactc gccgatgccatgcaggcggc agcacgtggagggggagggg 403 ageggggaettetteeegeg etgeetggeggateetggga tggtgageeetttaatgagg acteetgteecaatteetet 483 acggtccgtggatgccagga ggctatcccagctcgtggtc cgggcgtcctgcagagtgga acctccattggttccccgct 563 cccaattaagtaaaacgact ccacaggggtctgagtcgcc ggccttaggcgctccgccgg ccttaggcgccgcttggagt 643 tgctctctcccgttgctgtc ttgctggccatctcagcggc ctggcctccgccagtgtccc ggaggatgcagtggccatgg 723 ccaaacgccttttccataga ccctaattcaaaccagactg caggctgcacccccagcgcc gcggagtccgggcgctcggc 803 cctttgcaccggggcaagtt tgggcacagcagagccggcg cgggaacagggggaagctga cgttcggggtggcgggaggg 883 acgggattaaggctgtttgt gggacacaagagggtggctc agggacttcggtttttctct ggctgccccaggtgagccgg 963 gccgagctggcagcgggagg ttccgggaagttggcttcag aacgctgaagaccctaagaa cccaactttggggtcgctga 1043 agttgtgctgccccggagg gcctcctccgcatggcccgc gcgggggaccctccccgcga gtggaccccggtacggctct 1123 tecetteccegaetegget ttgtgetgaageegeggta gggaaggegggteeettgge eegeecagtagggeegeggg 1203 gaaagagggacgaacgtgga gctggcgactggtgggggaa gcttctgggtaggatgcagc catccacctttggtggggtc 1283 ggtctctctaatcagcggct tggcgacaaagagcttggtc gagggtaccccagaaagtgc tctcccgccccaagccgccg 1363 tegetageeegeetteecaa egggegetttgtteteggee eetgtaaceetteeetggga accgeeegeagegetggts 1443 cttgacgtgggccgggtcct gggtcgccgccagtgtcagc gctgccctccggtgtccacg cccctagcccccgcacccgc 1523 tgtgaagtcccgggtgtcct ttccactggcgctttgccca acccctggaaggcagaggcg aggtgcggagcctcaggctt 1603 tatecteceggaagtggeag teteceacegecacatetgg tetgettaacttegatagte etggeaaaggeagacaegtg 1683 cacagggaaggaggttgag cgctggtagataccaaggtc gtgtacaaataaagtggcac acgacacgtccccagtcact 1763 gttaatgcattgccttcgct ccttcccaggtggctggtgc tctccatcactctggagccc aagagagggcctccataatt 1843 gtattgcccatgagttgggg ttgtgtggggggcccaaatc agggttctctgggagggcta tgaattccgaactgagtctc 1923 ctgtgcactcctggctttaa ggttcaagaaattgtttgag ggttgtggtttttgtgggac tcagattatgcctggaatca 2003 tagttaccactgtggagaag aaagtggagctacttagcat gcctccccggcccgcctggc attacctccggctctgttct 2083 ctaggcccaacgtgaggcct cactggggcagtacagatgc agtactgaatttctttccag ccaggatctggagaggtggt 2163 gttctcttccctggtgtctt tagagaggcagatattcctg tgacctaagcccctcaagca cccattaataatgctgagta 2243 gacaactagaggtggcgttt tccggaacttcctgtgtgct ggcctgggaggttgaaccct ctaggaaacaggtctaggaa 2483 caaacaaacaaaaaaaacc ccacattgtttaaaagtggg tgcataagagtgaggacata ttcagagcttccccttttcc 2563 tgaaaaatgaaggcagctgg gatttacttaaaatgagagc acatatcacaattgccagag agctggtccctttctcaggg 2643 ctccctaagctcctgtggga agcaggtcagacagccctgg ggaccagagagatagggagt gcttttgggtgcctgccttt 2723 gaatggggaaggggggggg getgetgggatcagaggetg ctagcaactactccccagag actgaagcaggtttgtccct 2803 cagtgtcctgtggtcttctg tttctcctatatagaatagg agaaatggttatttgctctg gaatagtgacttgctatttg 2883 ttccctttctttcctctccc ttactgtaatcatttggact agtagagacactttccccag gtctggcagaatgggaggga 2963 gtgggggaggcctgtgcttg catgatgtcactgctggctt cagctctccagggagggtgg agttggttgtaacctacctg 3043 tggctcttgatgggccacaa taaaacctcattaacacaca ttggtagggagaagggactg gaaagaatgatgggaaagat 3283 aagtateettggttgttttt tgttgttgttgttgttgttgttt ttttgttttaagaca aggtttetetgtatagteet 3363 ggctgccctggaactcaata tgtagaccaggctggcctca aactcaaagaaatccaccta cttctaactttcagtgctgg 3443 gcctaaaggtgtaggccacc aaaagtgctcaacttttaca aagcagtcttactttgagca ggattctgaaacccttattt 3523 cctttctgttatcttcaaca atacactgctaggtgtattt agtccctcatgatgctgggc ctcctcaagtggcgccaggt 3603 caagcagteteetggttttt ggtggetetgaagaagaetg tgteecagtgaetggeagtt tgaatteggagettetettt 3683 teetteteagtetttggcag geagagtgaeactggtgtge eeaageetggagettetetg titaattetagtttatttte 3763 tttatcagactgaaaaacaa atcaggttggttataattct tataaacacgaaggtctcac ctttgcgtacgtctccggct 3843 gtgtgggtctgatgtccctc gggaatctctgttgaggctg ctgcagtgtgtgtgtgcgtgta gaaagggcaaggtagaatgg 3923 acagaagcgtgctgcccacc ccactgtcctgttcctaaat gatgaagcactggcccggtg aagagcctagagaactccct 4003 cggtgggagatgcacacaat gccaggaagcacacaggagc ttgagttccagcttggcagt gtcttctctttggtgacttt 4083 atcagetecagetgecetgg actaacaaacaaggetaget cacteteagtattgataate gaaggteettggttetgttt 4163 gagactgatcctcactcggt agccttgaactcttagcaat tctcctgtctcaactttcaa agagctgaaattacagactc 4243 gagccaccatatgcgactga aaccttgttcctaatccttg actgtgaacgactcttgggt ttggttctttctccatttct 4323 tragtgtatgttttagtrcg cgrcctacataatctattgc ccatacttagaaacaacagg tragagacagcattgggtcc 4403 agcagagcctcacactgaag ctcagtcctgccactgattt accgtgtcagctcaagtgac tcacttccaactcctctgct 4563 gtgcaaaagtgctttgtaag taaagtgctggggaaatgtt agctgtcgataatggttagg gttaactttttattgagtgc 4643 ctgttgtgtgtgggggttggg tggggttttttttagaggctt ggtagttttcttacttcttt cctacttagcttttcttcct 4723 aagcetttatggtatgtate attgeetgattgtttgagtg tgtgeactgaggeacgeetg tgeatgtttgagagtatget 4803 tgtgcgtgctctcgtgctca catatgtatggtgtgaatac actgtagagtgcaggccggc acactggggctggctgaatc 4883 ctgtgagccctgcctggagt ttgcagatcttccttggaca ctcctgcttgtgagcatttt gtgtggagtgactgtttagc 4963 tggctgtagcctacattgtg cetttgggtaaaccctgagt attgggaaacaccctgggct gtggctgtgtgtgcccgacg 5043 gttgcttgggtacagctaag aactcttcatagaaagttga gctcacatgctattagtatt aactgagtgctaaggaacct 5123 gtcttgggtggtacctgctt gccctctcatgcagtttatc ttgagcttggcgaacacact tacagatttagtagagcttt

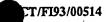
5203 tgtcagccctgggaggtggg tttcgtggccacaagtgggt agcttggaatccaagactcc tggcttctaggttgcattct



5283 cctgtggttctttccaaggg aatgctaggggaacattttg gacattagattatttctagt cccaaagcacacagaacata 5363 etgittectaattgeettit tittgittteeteteaatet ggitttgaagtgitgggitt gaaaattgeeeetgagage 5443 ctgccctagtgtgtgcagag ggaagatagtggaacaggaa gtctgtagaaagtatcttcc tttccaggaccttgtgcccc 5523 ggagcagagtcagcatggtg tcatatcgcttttggctatt ccagaagagatgaggtttta ggtgagaatgaaccttttag 5603 aaccttctagaaccttctgt tgagtatgacaggaatgccc tgaatagggtccgaagtgca tggccacttgtttgtctttt 5683 ccataagcaagcagcttcag gtacagacaataagactagg ttettggagtgagaccetgc acttggtgccatttcagctc 5763 cagatggacactggaggtcc ctacacagcaggctctggga tggctggctttgctatgtac tgttgcctgctctacaagag 5843 cttcccaggttactagcctt tgtcgacgctgggctcgctg gccaggcttgggcattggag aagggacaacttgccacctg 5923 gcataggctgtgtgtttgga gagtcaggaggtctggtgaa gcccgcaagtggaggcaagt ttagtgggacttgaggagag 6003 ctcagtaggaaatctctggg ctagtgacaggcaggtgtgg tggtggtggcgaggtggcgg gtctagatctccttttagag 6083 atttgcctagggatcgtccc tgctgactctggaactcaga ggcctccagaggtgtctcct ctgggagcctctcaagggtc 6163 teccatetectactgtttat ggetttgtgggetacetaat tacatagagaagatatgtte etetgeetecageeetggaa 6243 agttctgcccagtgactcac ctgagcctgcagccatgtgt gtacacaggcgctctcaggg gcttctgtcctgctggcttc 6323 agectttctageccetggtg ttctcggcagtggtagcatc tgggaaaccgggtcacctct tatttgcagctccctt 6403 tettggtgtettececettt ttaactactggtetgatgge ettagacteatgetgaaatt eteetttettttgteetage 6483 cttgtctctgacttcttgtg atcctctgggcctgtgaaat ccgctcaggggcctccattt ctaacagtcacacactggtg 6563 gagagaccgagtcctgggat ggtgaagctaaccctgctgg gcttctcaagcttcatttgg tttctctttattccttctgg 6643 aggtactgcctgcccaggg gagtctcagactagaccact ctggagttggaggtggggca ggttttcagatcagtgccct 6723 tggcattcgttgtgggaatg gggtggatggggcctctggg caaggtcaggctgggggtgg aggccaggtgatgttctccg 6803 cacccacacccaggcagcct ggcaccctccccaaggtccg ctcatcagcaggaatgaaag cagtgccgggcaggttgggg 6883 cagtgggcaggtgggcgtgt ttatcgctgtgctcatcagc tgagtcacgatgccaggccc cacaagtcctccctggaggc 6963 teacceaccaccttgace caccagcaccactagcagg aggtagggcagtgag acaagaccagcctgggggtc 7043 tgagaggcaaaggggagttg ttcatgacctggctgtgcat ggggacttgtgggtgtctca gatatctctgctgtccagga 7123 ggaagetgtettaagtgeea acetgeetagageeeetget gggtgeaggaaatgeaeaag ggagagtgeeeateeatgga 7203 ataggcccatggagctagac cagtgacagtgacagtgaag tcagcccccacctgtgtctt ccgagccagctggagggttt 7283 tratctcagattctgcgaaa ccatagaatctagtcaggag cctagactgcaaagcaggct tcgttgatgctttaacttgc 7363 aggetteetgggtatgaggg ataettagaaaggteeegea ggtagggagggeateaggaa gtagaagagggeeaggeact 7443 totatotootgcattgcccc ottotoccatctccaaggat ggtaaaaagaaccettccag tacactgacagagagaaaa 7523 cccttcatctcaccccattt ggatctgtcgtatcagcatg tgctggccctgcttccatac cagaggtggctagagatgtt 7603 ccctgggaattcactggttg gggacttgagtgtatcagag gggcacaaagtaacattaac tctggtatcctctgcagcaa 7683 atcggagatcccctctccta ggcgagttctcagtggatat ggaggtcaggtttgggcttg tagggccccagcaagagtcg 7763 ttgatgtcactccagcttct cccgaggaagatgagggtgc tgtgttgggatcacatctct ccctgaatggcatgttgggg 7843 agggatggagccettgette tgacccctaagcttggtett taggtggccacagtetetgg gttetgteetaceteeetge 7923 ccttgtgtgcttcaaaggca tgctaaagggactctcggcc attccgaatggcacagtgtt ccttctgttctcccaccccc 8003 agaaggaggcaggcctggat tgtagattcctagaagtaag tggccctgagcatgctgttg atgaacctggaaccaggcag 8083 gctgggcatcctaggacctg tctttccatagaagtctgaa tcagtctacctttgggactg agtaaggggctcctcacata 8163 tcagctggctagtccatctt ggctgatctaaaccacatta ggctgaagagaagcatggtg tacagtctggtccacccgaa 8243 ccacatactggctttatcag ttctcgtataattttgcagg taactttttagctctaagcc tgtctcctcatctgtgaaat 8323 egggteeeteatateetgee tagaagggettttgaaaaga ttaatgaagtagtatgeega gtggttggggtteteteteett 8403 gactggagcaagtctctagg agtactaaggatagcctgct gtgtgcagcacccccaggga ctgtgcctgagtaggagggt 8483 acagagtetteatgtgaatg gecettetggtettgeeeg aagttagtgttgatgteata gagtetacaaacatgeettt 8563 tgtccttcctcagaagtcca agcctttcctggcagaccag acattcatctccactgagcc tctatgtgagactggctcct 8643 ggcctgagctgtgtgggctg agctggcgaatgggaaaact agacacctgggcacctgggt gggggctcgggacagcagtg 8723 tttcagttgtaggcactgtg cccctgcctggagcttctga ctgaaggttaccctgagagg aagcaggttccctatagaca 8803 ctaacatagetgggtcagag tgcaaggtgggtgtgcccct gccctgacccattcagtgca aaggctgctcttctgggagt 8883 gagagetetgacaggaetgt gatggeegaggggteteaga geaaacetgeetggeetete eccaetetgatggatatgtg 8963 ctcttaaacaagtgactgtc cactttgcctcaatttcaac atctgtaagatagatagggc gttatggtctgaaaatggtt 9043 ttaaagattagttagetaat acagggaaagtgetetgaca ggtacetggcacettaetea acaagtggetggagtgeetg 9123 atttectaaggtetegacet gtecetatgetteaagtgee cetacageettggteaggee ettaggtteteceaeceaec 9203 gctggccccaggacctagac tgctggaccctgaccccatt tttcctttaagccacctctg cgtcaactctaaaaggcggt 9283 ggagttgtttatctaggctg tgaggtgtcagagaaaggac ctgggccgctttgttcctgt gtgggctggggccactccag 9363 gaactgagaaacccaccac cttttcaaaaacagcctctt ctcagagtctggcacctcag ctagccaccatgctgtggga 9443 ccacteccageatgetetge etttggtttgttteccaggg geeteagtgeettttaaaga tgcacaggeatetttagtte 9523 aaggggaaagaggaaatgaa gtgtatttgctggtggtggt attcctgtcacttgcattct cacagaggctaaagaaattt 9603 gctctttgtatcttctagtc tcttctttatgatctttcc catctgttgtatcccaactg cagggccccagttctagaat 9683 tagecceteccecataggaa geegaettatgetataatgt gaatgacaagtateetttag eeetteecacaggeatttta 9763 attttcaaaagggcattgca caaccgcagagacactaaga agagaggtttggtgatcaga gttacagccccagcctccca 9843 gctggtggcccggctggtgc aggtgtgtcgaaagcagtag tttctgcttcagtgaaactt gaggatcctttatttagcca 9923 gttcaggggcggaatggcca tgcgaagtctatgtgtcaca ggtgtcaggcccccatatcc tgctgagtctagaatcagct 10003 acgtagcagttttgggggta ttgccagactgggagtttac atcccagaagcgagaatggt ggggttcctatactgctcca 10083 gacaggatetttcccccaag tttgtcagccacctctcttc aagtcccttggctctgacca gcaagacgtatccaaaagaa 10163 actgaggaggcccttcactt ctttttaggatagtgtgggg ccagcatggtgggggttggg aatggctttctgtctcttcc 10243 atcacaggctacttccc agagacactttgattctggg catctccagcagtcacctgg cccacaatgctttgctgccc 10323 tttgcttcagccactgtatc tggttgtcccttgaaggtga gccagagctcctaggcagag agcatgtgctatacaaagcc 10403 gtaggetgggecetgggaac cttcttgetgtcatcctcct gtcaaacccctatggtatgg tagcccacataaggcttgtg



15603 gggtgctcttctccgccgct gaaggaggcagccgcaggga agactaccacaggaatccga gtaccacctggagcagtgta



15683	tacaggatgtgggctgatgt	gtggtaagggcatgatgggc	tgatgtgtggtaagggcatg	ggatctgattgctctgtgga
15763	tigggcgacagggacattttt	gagtgtctactgcagtagtt	ctcaacctgtgggttgtgcg	ccccttggtgggagttacat
15/03	- trace tattace trace	ttcataactgtagcaaaatt	acaattotoaaagaaccaag	aaatcaccgcagcatgagaa
15043	actagatactacactacga	tgttaggagggttgagagcc	acteatectetagatetaga	ccatggcgggctgtaactgc
15923	cctgtattaaagggtcacgg	tgaaccagctgtccttgcag.	at an act to to a source con	aacctttgtcccagggaga
16003	tctctggagttaagccacag	tgaaccagctgttcttgtag.	act at the acceptage to the	actatatataasataccc
16083	agagettgettttgetttgt	acttttaaaggaagttcagt	ggtettegggeettgtgget	getgegegeggaagegeeee
16163	tgtacaataagctgtataga	tcgtgtacaactgcagtttt	cctccgtgggtccaccaacc	actectgactecacggatga
16243	gtgaggccagtagggctgtg	tgtgggtccctaggccaagc	atcctggaccacgatgagcc	tcagctagaccactctggat
16323	ctttagcagaggctcctaga	gagetggetgetteeteet	gccttcttttctcttaaaac	ttcgtctcaatcggaagctc
16403	ctctgtgcacgtgacctcca	ggcctgggggtcgccacaaa	tccctcatcacaagacgag	cagctcgcatgagggacacg
16483	acacttottacctaccago	tgtggggtttttgttggttg	gttgttttgttttgt	ttttttacttgtacagaagt
16563	attataacatcaaatatcaa	ctgttagtgctggcaccatt	ttacaggtagggaactgagg	ctgtaagatgtgtagtgaca
16643	teactanaceacteaatta	gtgaggccttaccaaggtca	ggtctttggagccttttgct	gaaccatgtacttctatctc
16722	tegetdaggetaeteagetg	tatatggctctggctagcct	ataacccatatgtagacga	ggctgacctcgaatacactg
10/23	tgttttgttgaaacaaagtc	ctgggtggcaggattgaagg	catchesttectectaacte	tacactttaaaaaaaaaaatc
16803	cagtettttatgtetgeett	Cegggeggeaggaregaagg	catgigatic cetectaceg	tagattagtctcattactga
16883	attetttgttetggtetgtg	ccagggccttgtaagatgtt	ccgcgccgagccgggccacc	
16963	gcagggcccctgtatcttcc	ttctctgtcacttgcttacc	tgggtetteeteetgeacta	getatectagaactagtact
17043	gagagcaactatgggcccaa	ctctgccccttgcccagcct	gcttagctgggggcggtgtt	ceaettecetgeceaagtee
17123	tgtgggactgtgtttgtact	ccaccaccttcagttccttg	gagctggagcaggccaggcg	gctgcattcctgcagctgct
17203	gttgccagggagagcccatc	ccattcacttcagtctcctt	aatgtagaagccttgtcgaa	ttagcttccactgtccccaa
17283	cccaagagtaccctgtcctt	tcttcactaagaaggccagg	atacagtccttcctgtggct	gataagacaggccttgggac
17363	aaggctgggaccacactgt	gtgggcaaagctgcttcagc	accgatggctcctccatgcc	aagcttggctctgcttctca
17443	cagttgagacttctgtgcgc	acaccactgtctagctcag	ctggacactgattttcttta	aatgtatagattttggggtg
17522	aggrafactassagctcccs	ctgatgccccaagcctgagt	ctcagagtatgatcaattga	togctttcatoggtatcaca
17502	gggtgtgtgtgaaageteeta	ctccctgaccagtcagagca	tectagggttagacaatgtc	eccatcacttatacctccac
17603	gereegereeggeraga	tatggcattgagggtatgag	aagaccagggtttcccag	agttacgccaggcgcacag
1/683	etggcaccaggctatgatgt	tatggcattgagggtatgag	aaggaccaggggcccccag	taggetaggetagg
1//63	gcaattgtttcctacatgtg	tggctggaatggttgggtga	at an act at a support of the control of the contro	A A A TOTAL CONCOUNT A A A TOTAL CONTROL OF THE A A A TOTAL CONTROL OF THE A A C A TOTAL CONTROL OF THE A C A TOTAL CONTROL
17843	gcattgaccaaggcatatct	catacccttttcttatcttt	Classical	AAATGTTCCTCCTGAAGATC
	•		GINIIevalAlava	lAsnValProProGluAspG
18923	ACCATECETETECCCATEAC	TCTGACAACTTCTCTGGCTC	TGGCACAGgtaagactgacc	cagaacactgagatggcata
10,23		SerAspAsnPheSerGlySe		
	IIIASpolySelolyAspasp	BEIMSPASMI MEDEL CLY DE		
18003	gatcatggctggagtggtga	gcaggcagtcacccagcttt	tagtgaacccccttcttctc	ccatcccatccttagccatt
18083	ggagtcaggacagtgccaaa	aggaagaatggtatccagct	gcaagccactcagctaagag	aaactctcagagaaatgaag
		cagccactagagccaaccct		
18243		tttgcccaagagcctgttag		
		accagttctctgcagagctt		
		gggtctcccagaagaatggg		
		ggtaggtaaaatgcagccgt		
		cagtcctgtatcagctctgg		
		cagagagcacgctacttgta		=
		tgctaggtcacctttggggt		
18803	aggcgtgatgctggagacct	ctcctaccctgacaccccaa	accccacctctgaccctgc	
	•			lyAlaLeuProAspThrL
10002	momes cooks as as commen	3 CMMCC3 3 CC3 CCMCMCCCM	CMMC) C) CCC) CCCC) C) C	COCC 1 C C C C C C C C C C C C C C C C C
10003		ACTTGGAAGGACGTGTGGCT		
	euserargGinThrProser	ThrTrpLysAspValTrpLe	uLeuThrAlaThrProThrA	laproGluproIniserser
19963	AACACCGAGACTGCTTTTTAC	CTCTGTCCTGCCAGCCGGAG	AGAAGCCCGAGGAGGGAGAG	CCTCTCCTCCATCTAGAAGC
17703		rSerValLeuProAlaGlyG		
	ASHIMIGIUIMATAFNEIN	15el valbeurloalagly	1 day serogradia day dia	Plovalbeunisvalgiumi
19043	AGAGCCTGGCTTCACTGCTC	GGGACAAGGAAAAGGAGGTC	ACCACCAGGCCCAGGGAGAC	CGTGCAGCTCCCCATCACCC
	aGluProGlvPheThrAlaA	rgAspLysGluLysGluVal	ThrThrAraProAraGluTh	rValGlnLeuProIteThrG
	_			
19123		GTCACCACAGCCCAGGCAGC		
	lnArgAlaSerThrValArg	ValThrThrAlaGlnAlaAl	aValThrSerHisProHisG	lyGlyMetGlnProGlyLeu
19203		AGCACCTGGTCAACCTGACC		
	HisGluThrSerAlaProTh	rAlaProGlyGlnProAspH	isGlnProProArgValGlu	GlyGlyGlyThrSerValIl
10292	CAAACACCMMCMCCACCAMC	GAACTGCCAATCAGCTTCCC		1011
19203				
	enyseiuvaivaiGiuAspG	lyThrAlaAsnGlnLeuPro	AladiyGluGlySerGlyGl	uGIN
19363	ttectaggtagecactagta	cctgcacctggccgcctaat	atcctcattacagtgacego	taacaaaateeeacetteet
19443	CCtdccdaaacadactdgcg	tgcaagatcaggaggtgggc	gactccttagatgtc=ttc=	duanttacanceanctae
19522	ttttccgtcttscscstsc	tgggaattttcacacaacaa	tatatacattatatata-	agagatatatatatat
10602	cotagesageteea	gactgactgcattccttgac	antactactactactactactactactactactactactact	aygeyyeardegegeeergg
19003	cergeeagggreecaggrgt	gattgattgeattgettgac	ayatyetygtataggttggd	Lacgicigatgggggtggca

19683	ggggatcccatcaggtatgg	cactgctcaggttgctgttg	tgtcagtggctccagctgac	ctgatcccaacctacccttc
19763	tgtagGACTTCACCTTTGAA	ACATCTGGGGAGAACACAGC	TGTGGCTGCCGTAGAGCCCG	GCCTGCGGAATCAGCCCCCG
			aValAlaAlaValGluProG	
	,			
19843	GTGGACGAAGGAGCCACAGG	TGCTTCTCAGAGCCTTTTGG-	ACAGGAAGGAAGTGCTGGGA	Ggtgagtcttctttcaggtg
	ValAspGluGlyAlaThrGl	yAlaŞerGlnSerLeuLeuA	spArgLysGluValLeuGly	G.
	- -			
			tgctggggtgaagcatcttt	
			gtctagtctgggagagacta	
20083	ctgatctgagcctgcctctg		GCCGGAGGCCTAGTGGGCCT	
		lyValile	AlaGlyGlyLeuValGlyLe	uIlePheAlaValCysLeuV
20162	maccommes macmams acco	10011011011001001100	ca coma omo cimpo ca coa co	CC111C11CCC11FCCCCC
20163			CAGCTACTCCTTGGAGGAGC	
	alalaPheMetLeulyrArg	MethyshyshysAspGluGI	ySerTyrSerLeuGluGluP	rolysGinAlaAsnGiyGiy
20243	CCCTACCAGAÁACCCACCAA	CCACCACCACTTCTACCCCT	GATGGGGAAATAGTTCTTTC	TCCCCCACAGCCCCTGCCA
20247	AlaTyrGlnLysProThrLy		0.110000.4811.101101110	100000000000000000000000000000000000000
	Alarytonnbyseronniby	Sollioluolurnelylytä		
20323	CTCACTAGGCTCCCACTTGC	CTCTTCTGTGAAAAACTTCA	AGCCCTGGCCTCCCCACCAC	TGGGTCATGTCCTCTGCACC
20403	CAGGCCCTTCCAGCTGTTCC	TGCCCGAGCGGTCCCAGGGT	GTGCTGGGAACTGATTCCCC	TCCTTTGACTTCTGCCTAGA
20483	AGCTTGGGTGCAAAGGGTTT	CTTGCATCTGATCTTTCTAC	CACAACCACACCTGTTGTCC	ACTCTTCTGACTTGGTTTCT
20563	CCAAATGGGAGGAGACCCAG	CTCTGGACAGAAAGGGGACC	CGACTCTTTGGACCTAGATG	GCCTATTGCGGCTGGAGGAT
20643	CCTGAGGACAGGAGAGGGGC	TTCGGCTGACCAGCCATAGC	ACTTACCCATAGAGACCGCT	AGGTTGGCCGTGCTGTGGTG
			GTGGGGAGGTCTACTTTAGA	
			TACTGGGGTAAGGCAAGTCT	
			TAATAAGACTTTTTTTTTTTT	
			GTGTGTTGTCTCTGAGTTTG	
			TAAGTAGGCTGGCTAGTCCA	
			CGGCCTTGGCCTGGGCCACC	
			GAGGAGACTGTAGAGTTTTG	
			TTCCTCCCGGAAGTTGACAA	•
			TCCTGTGGCCCGCCCCCAG	
			ATCCAATGGGTTCTGTACAG	
			GCTTCTTCTTCTCTGGACTG	
			GGCTGCACGCCTCCAATCCC	
			GTGTTGTGTTTTTTTTGG	
			CTTCTAAATGGTTTAAGTAA	
			ACACCTGGGACAGCCTTCCT	
			AGCTCCTGGCTAGGCTGCTC	
			TCACATGGCTGGATGTGATG	- -
			CAAGCCTCAGGCTGGCGCTC	
			tgtgcagtctctttttggtc	
			cttggtaccaaatagcaaca	
			cccgactttgcacctgtttt	
			attctgctgactctgaatgt	
			ctcatgtgtcaggattacaa	
		•		

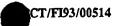


Fig 4.

TCTAGAACAC	TTATTAAGAG	CCAGGCACTG	AAAAGTGCAG	ACTCCCTCAT	TTCATCCTGG	60
CCGTGCTTAC	AAGTAGTTTC	CATGCTCTGG	TAACCCTGTG	CAGAGGGCAG	CGTGGGAGGC	120
GGGCCGCTTG	GTGGACGGTC	ATGGGGGCTC	TGCATGGGTG	GTTGCCCTTG	CCTCAGAAGA	180
ACTCCCTAAG	TAAGAGCAAG	TTAGCCTCCC	TAACCCCTGG	TGGGTTGTTG	CTTCTTTTCT	240
CCTCTTGTTT	CTGCCAAGAG	AGGGTGGACC	AAGAAGACCC	CAGCCTACAG	AACATGTGAT	300
CCAAATAAAC	TTCTTTTTAG	TATAAATGTC	CTAGCCTGTG	ACGTTCTGGT	AGACTAGCAC	360
AAGATGGACC	AAGACAACTC	TCATCGAGAC	TCTGAGGAAC	GAACTGGCAT	CACATGGGAA	420
CAGGAAATGA	AGCTTAGAGA	GAGGTTCTGT	GGCTTGTCCA	ACATGGCTGT	AGTTTAAATC	. 480
CAGCTTGCCA	CCAAAGCACA	CACATTTCAC	TGCTGTGCTG	GGCCGGGCCT	CAGATCCCAG	540
GGGCTCCGGA	GCTAGAAGGA	CACGTGTATC	AGCCATGGCT	TCAGTTTATT	GCTGTATACT	600
CTGTGCTTCT	GGCTCTCATG	GAAAAGACAG	ACATTGGGGT	TCTTATAATC	TCTCCCTCTC	660
CCCTCCCCAC	ACTCTATCCC	CAAAGGAGGC	ACCACTTCTG	CAGGTAAATG	TTATCTTCAA	720
AGCGCTCACA	TCGCAACCTT	TGCCCACACC	ATCTCATTAA	AGGAATTGGC	AGTGACTTTA	780
AGGTGAAAGA	ACTCGGTGGC	TACGTGTTAT	ATAAATTTGC	ATCTGGGTCT	CAGAGCTGGA	840
AGGAAGGCAC	TCCCATACAT	GCAGTCTGTA	CATGCAGTCG	GATGATGGAC	CAACAACACA	900
TTGTGATTTA	TGCCCCTGCT	GGTGAGCCCA	GGAATCCCTG	TAGCACTCTC	TCTCAGCTCT	960
AGGGCCCTGC	TTGTGTATGG	AAAACGCTTA	GTGTTTTATA	GGTATTTTGT	CAGAATACTT	1020
TAAGGAACTT	GACCAAAGTT	ACAGGGAGGT	TAGACAGATT	GTCATGGTAT	ACTCACCTCT	1080
GTCTCTGACC	CTCCTAACTG	GGACCTCTTT	AGTCTCCCTT	GAGGCAGGGA	GTGCCACATG	1140
CATGTGTCCA	GGCACATGTC	TCCTGGTTTA	CCTCCCAACG	CACCTCAAGT	CCCCAAGGTA	1200
GGTAGGCACT	TGTATTCTGT	AATTCAGAGA	GGCAAATCAA	ACTGTTACAA	TGTTTGCCCA	1260
AAGCTCCCCA	AGCAAAGTGG	CCCTAAGAGT	GAGCAAAGAG	ACTGCGTGCC	TTCACTGCCT	1320
GTGTGAATCC	CTGCAGATAG	TCTCTCATCT	TGGTGCCCTT	CCCACAGAGG	CTGGGGCGGC	1380
AGGAGGGAGC	CTGGACAGCT	CAGACACTGG	GTCATTGATG	ACTGTTGTGT	GGGATACCTG	1440
CCGGGGCGCA	GGAGTGAGCC	ATGCCACCCC	AGGAAGTGGT	TCAGGGTGAC	TCTTCTTGGC	1500
ACACCTGGGA	GGATGTAGCT	GGTGCTGGCA	CACCCACCGT	CACGAGAGCT	TCCTGTCCAA	1560
ACCTTCAACA	AAGGCGGCTT	CTTGAGACAG	GCTAGACTGA	AGTCACCAGC	CTTGGGTGGG	1620
GTCCACTATG	TAACCTCAGT	GCTCAGGAAC	CCTTTCCCAT	ACTGTCTGGA	ACTATACTGT	1680
ATGTAGCTGG	GTTTCCACGC	ATGTGTGCCT	GCACCCAGTC	CATCTCATCT	TCTATCTCCC	1740
TCCCCTTTCC	CGCTTCCCCC	CTCCCCACTC	TCCATCTCAT	CTTCCATCCC	CACCTCTTCT	1800
					ATGTTCCAGG	
CTTCAGTCCC	TTCTCTGCAC	CTGTCCTTTG	TGAAGTGACC	AGAGGATTTC	TGATCCTGTC	1920
TCTGTCGCTC	TGAAGGGTCA	GGAGTTCCTC	CTGCCTGGAC	AAAGCCATCC	TGACGCACAT	1980
					TACTTACAGG	2040
			GGGGGGGG			2100
				AATAAGGGAA	AATTATTCTG	2160
GATTATTTGA	GTGAAGCTGA	AAGGTGATCA	TCTAGA			2196

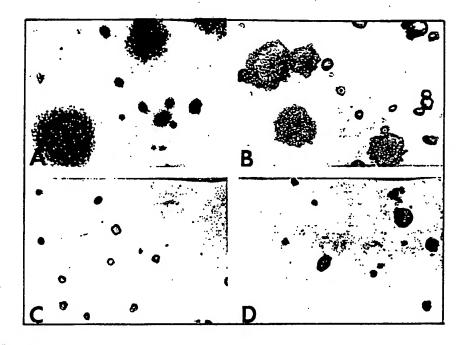
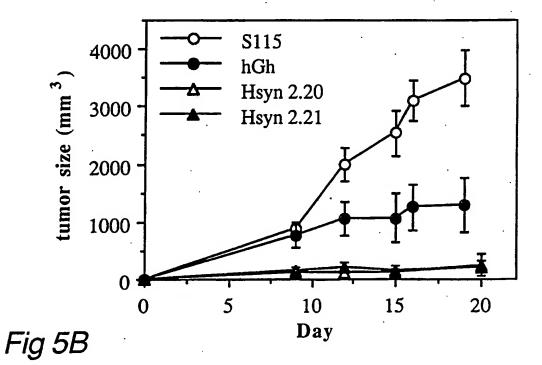


Fig 5A



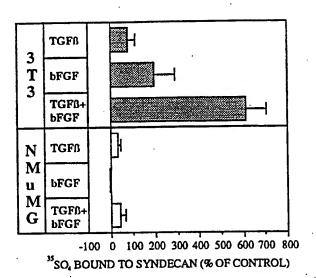


Fig 6.

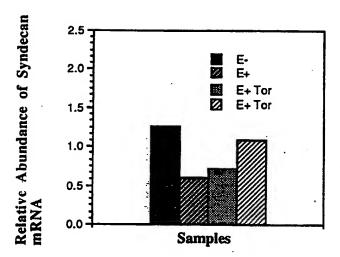
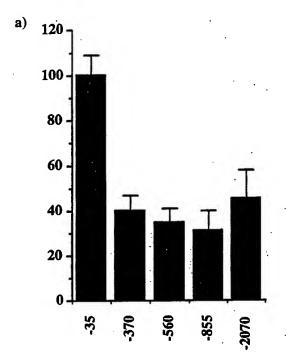


Fig 7.



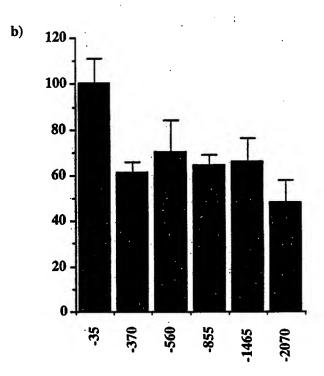


Fig 8.

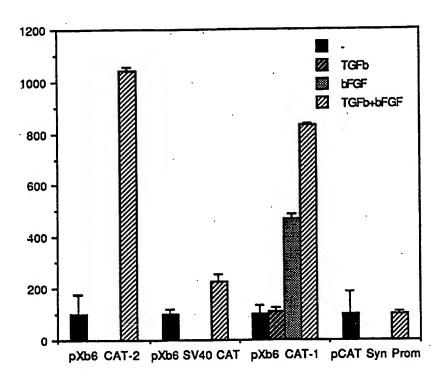


Fig 9.

mal Application No Inter 3/00514 PCT/

A. CLASSIFICATION OF SUBJECT MATTER
IPC 5 A61K31/135 A61K37/02

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

 $\begin{array}{ll} \text{Minimum documentation searched (classification system followed by classification symbols)} \\ \text{IPC 5} & \text{A61K} \end{array}$

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

Category *	MENTS CONSIDERED TO BE RELEVANT Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	CELL REGULATION vol. 2 , 1991 pages 1 - 11 S. LEPPA ET AL. 'STEROID-INDUCED EPITHELIAL-FIBROBLASTIC CONVERSION ASSOCIATED WITH SYNDECAN SUPPRESSION IN S115 MOUSE MAMMARY TUMOR CELLS' cited in the application see the whole document	1,3-7,9, 10,13, 15-19, 23,24, 26-32
Y	PROC. NATL. ACAD. SCI. vol. 89, no. 3 , February 1992 pages 932 - 936 S. LEPPA ET AL. 'SYNDECAN EXPRESSION REGULATES CELL MORPHOLOGY AND GROWTH OF MOUSE MAMMARY EPITHELIAL TUMOR CELLS' cited in the application see the whole document	1-6, 9-18, 21-31, 34,35

X Further documents are listed in the continuation of box C.	X Patent family members are tisted in annex.
*Special categories of cited documents: 'A' document defining the general state of the art which is not considered to be of particular relevance 'E' earlier document but published on or after the international filing date 'L' document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) 'O' document referring to an oral disclosure, use, exhibition or other means 'P' document published prior to the international filing date but later than the priority date claimed	"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art. "&" document member of the same patent family
Date of the actual completion of the international search	Date of mailing of the international search report
1 March 1994	2 8. 03. 94
Name and mailing address of the ISA European Patent Office, P.B. 5818 Patentiaan 2 NL 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax (+31-70) 340-3016	Authorized officer Hoff, P

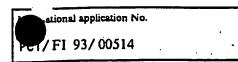
Inte	mal Application No		
	/FI	93/00514	

	tion) DOCUMENTS CONSIDERED TO BE RELEVANT	Relevant to claim No.
Category *	Citation of document, with indication, where appropriate, of the relevant passages	
Υ .	BIOCHEMICAL SOCIETY TRANSACTIONS vol. 19, no. 4 , 1991 pages 1069 - 1072 M. JALKANEN ET AL. 'SYNDECAN, A REGULATOR OF CELL BEHAVIOUR, IS LOST IN MALIGNANT TRANSFORMATION'	1-6, 9-18, 21-31, 34,35
• .	see the whole document	
Y	WO,A,92 13274 (JALKANEN) 6 August 1992	1-6, 9-18, 21-31, 34,35
•	see abstract see page 7, line 15 - page 13, line 3	
Y .	JOURNAL OF CELLULAR BIOCHEMISTRY vol. SUPPL., no. 15F , 1991 page 223 M. JALKANEN ET AL. 'STIMULATION OF SYNDECAN GENE EXPRESSION IN MESENCHYMAL CELLS BY bFGF AND TGF-beta'	1-6, 9-18, 21-31, 34,35
A	see the whole document	36-41
Y	THE JOURNAL OF BIOLOGICAL CHEMISTRY vol. 267, no. 9 , March 1992 pages 6435 - 6441 K. ELENIUS ET AL. 'GROWTH FACTORS INDUCE 3T3 CELLS TO EXPRESS bFGF-BINDING SYNDECAN'	1-6, 9-18, 21-31, 34,35
A	cited in the application see the whole document	36-41
X .	EP,A,O 335 554 (UNILEVER) 4 October 1989 see abstract see page 3, line 40 - page 5, line 26 see page 18, line 39 - page 19, line 40; claims; example 2	36-41
X	EP,A,O 455 422 (MERCK & CO.) 6 November 1991 see abstract see page 7, line 12 - line 27; claims	36-41
A	THE JOURNAL OF CELL BIOLOGY vol. 114, no. 3 , 1991 pages 585 - 595 K. ELENIUS ET AL. 'INDUCED EXPRESSION OF SYNDECAN IN HEALING WOUNDS' see abstract see page 590	36-41
	-/	

Inter. nal Application No
PCT/1993/00514

	·	PCT/F 3/00514
C.(Continua	non) DOCUMENTS CONSIDERED TO BE RELEVANT	Relevant to claim No.
Category *	Citation of document, with indication, where appropriate, of the relevant passages	Rejevant in claim 1405
A	J.E.F. REYNOLDS 'MARTINDALE, THE EXTRA PHARMACOPOEIA' 1989 , THE PHARMACEUTICAL PRESS , LONDON see page 650 - page 651 see page 1625	1-23
A .	EP,A,O 462 398 (HOFFMANN-LA ROCHE) 27 December 1991 see abstract see column 1, line 52 - column 2, line 12; claims; example 2	1-23
A	THE JOURNAL OF CLINICAL INVESTIGATION vol. 80, no. 5, 1987 pages 1516 - 1520 L. SCHWEIGERER ET AL. 'BASIC FIBROBLAST GROWTH FACTOR AS A GROWTH INHIBITOR FOR CULTURED HUMAN TUMOR CELLS' see the whole document	1-23
		·
٠		





Box I	Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)				
This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:					
1. X	Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely: Although claims 1-41 are directed to a method of treatment of the human/				
<u> </u>	animal body the search has been carried out and based on the alleged effect s of the compound/composition.				
2. X	Claims Nos.: because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically: please see attached sheet/				
•					
3.	Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).				
Box II	Observations where unity of invention is lacking (Continuation of item 2 of first sheet)				
This International Searching Authority found multiple inventions in this international application, as follows:					
1.	As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.				
2.	As all searchable claims could be searches without effort justifying an additional fee, this Authority did not invite payment of any additional fee.				
3	As only some of the required additional search fees were timely paid by the applicant, this international search report				
	covers only those claims for which fees were paid, specifically claims Nos.:				
4.	No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:				
Remark	on Protest The additional search fees were accompanied by the applicant's protest. No protest accompanied the payment of additional search fees.				
	140 protest accompanies are payment or acamania sea on term				

International Application No. PCT/FI 93/00514

FURTHER INFORMATION CONTINUED FROM PCT/ISA/210

2. OBSCURITIES,...

A compound cannot be sufficiently characterised by its pharmacological profile on its mode of action as it is done by the expression like "compound that induces expression of syndecan". Therefore the search was limited to the compounds mentioned in the claims 8,12,20,22,33,35,41.

INCOMPLETE SEARCH: CLAIMS SEARCHED COMPLETELY: 8; 12; 20; 22; 33; 35; 41 INCOMPLETELY: 1-7; 9-11; 13-19; 21; 23-32; 34; 36-40

information on patent family members

Inten 121 Application No

/FI 93/00514

Publication

Patent document cited in search report	Publication date	Patent family member(s)		Publication date
WO-A-9213274	06-08-92	AU-A- EP-A-	9071391 0569367	27-08-92 18-11-93
EP-A-0335554	04-10-89	CA-A- JP-A-	1325598 1287013	28-12-93 17-11-89
A Thomas Contraction of the		JP-C- JP-B- US-A-	1770969 4060567 5037643	30-06-93 28-09-92 06-08-91
EP-A-0455422	06-11-91	JP-A-	4224522	13-08-92
EP-A-0462398	27-12-91	US-A- AU-B- AU-A- CA-A- JP-A-	5147854 636489 7713991 2042973 5213772	15-09-92 29-04-93 12-12-91 23-11-91 24-08-93

THIS PAGE BLANK (USPTO)

This Page is Inserted by IFW Indexing and Scanning Operations and is not part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images include but are not limited to the items checked:

□ BLACK BORDERS
MAGE CUT OFF AT TOP, BOTTOM OR SIDES
☐ FADED TEXT OR DRAWING
BLURRED OR ILLEGIBLE TEXT OR DRAWING
□ skewed/slanted images
COLOR OR BLACK AND WHITE PHOTOGRAPHS
☐ GRAY SCALE DOCUMENTS
☐ LINES OR MARKS ON ORIGINAL DOCUMENT
☐ REFERENCE(S) OR EXHIBIT(S) SUBMITTED ARE POOR QUALITY
□ other:

IMAGES ARE BEST AVAILABLE COPY.

As rescanning these documents will not correct the image problems checked, please do not report these problems to the IFW Image Problem Mailbox.

THIS PAGE BLANK (USPTO)